Mikael Hartman

List of Publications by Year
in descending order

193
papers	12,008
citations	41258
49
h-index	31759
101
g-index
206
all docs	206
docs citations	206
times ranked	17453
citing authors

#	Article	IF	Citations
1	Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94.	13.7	1,099
2	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. Nature Genetics, 2013, 45, 353-361.	9.4	960
3	Menarche, menopause, and breast cancer risk: individual participant meta-analysis, including 118â€˜964 women with breast cancer from 117 epidemiological studies. Lancet Oncology, The, 2012, 13, 1141-1151.	5.1	753
4	Familial Risk and Heritability of Cancer Among Twins in Nordic Countries. JAMA - Journal of the American Medical Association, 2016, 315, 68.	3.8	648
5	Breast Cancer Risk Genes â€" Association Analysis in More than 113,000 Women. New England Journal of Medicine, 2021, 384, 428-439.	13.9	532
6	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. Nature Genetics, 2015, 47, 373-380.	9.4	513
7	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. Nature Genetics, 2013, 45, 371-384.	9.4	493
8	Genome-wide association studies identify four ER negativeâ€"specific breast cancer risk loci. Nature Genetics, 2013, 45, 392-398.	9.4	374
9	Mammographic Density Phenotypes and Risk of Breast Cancer: A Meta-analysis. Journal of the National Cancer Institute, 2014, 106, .	3.0	261
10	Functional Variants at the 11q13 Risk Locus for Breast Cancer Regulate Cyclin D1 Expression through Long-Range Enhancers. American Journal of Human Genetics, 2013, 92, 489-503.	2.6	201
11	Incidence and Prognosis of Synchronous and Metachronous Bilateral Breast Cancer. Journal of Clinical Oncology, 2007, 25, 4210-4216.	0.8	182
12	Incidence and Outcome of Male Breast Cancer: An International Population-Based Study. Journal of Clinical Oncology, 2011, 29, 4381-4386.	0.8	175
13	Exome sequencing identifies highly recurrent MED12 somatic mutations in breast fibroadenoma. Nature Genetics, 2014, 46, 877-880.	9.4	172
14	The Heritability of Prostate Cancer in the Nordic Twin Study of Cancer. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 2303-2310.	1.1	169
15	Genomic landscapes of breast fibroepithelial tumors. Nature Genetics, 2015, 47, 1341-1345.	9.4	167
16	Triple negative breast cancer in Asia: An insiderâ€™s view. Cancer Treatment Reviews, 2018, 62, 29-38.	3.4	148
17	Longer Examination Time Improves Detection of Gastric Cancer During Diagnostic Upper Gastrointestinal Endoscopy. Clinical Gastroenterology and Hepatology, 2015, 13, 480-487.e2.	2.4	136
18	Genome-wide association analysis in East Asians identifies breast cancer susceptibility loci at 1q32.1, 5q14.3 and 15q26.1. Nature Genetics, 2014, 46, 886-890.	9.4	135

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19	Risk of death from cardiovascular disease following breast cancer: a systematic review. Breast Cancer Research and Treatment, 2017, 164, 537-555.	1.1	132
20	Female Breast Cancer Incidence Among Asian and Western Populations: More Similar Than Expected. Journal of the National Cancer Institute, 2015, 107, .	3.0	127
21	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. Nature Genetics, 2016, 48, 374-386.	9.4	125
22	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. Nature Genetics, 2020, 52, 56-73.	9.4	120
23	Breast cancer in a multi-ethnic Asian setting: Results from the Singapore–Malaysia hospital-based breast cancer registry. Breast, 2011, 20, S75-S80.	0.9	115
24	Mammographic density and ageing: A collaborative pooled analysis of cross-sectional data from 22 countries worldwide. PLoS Medicine, 2017, 14, e1002335.	3.9	108
25	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. Nature Communications, 2014, 5, 4999.	5.8	105
26	Fine-Scale Mapping of the FGFR2 Breast Cancer Risk Locus: Putative Functional Variants Differentially Bind FOXA1 and E2F1. American Journal of Human Genetics, 2013, 93, 1046-1060.	2.6	98
27	Breast cancer research in Asia: Adopt or adapt Western knowledge?. European Journal of Cancer, 2013, 49, 703-709.	1.3	98
28	Simvastatin-induced breast cancer cell death and deactivation of PI3K/Akt and MAPK/ERK signalling are reversed by metabolic products of the mevalonate pathway. Oncotarget, 2016, 7, 2532-2544.	0.8	95
29	No evidence that protein truncating variants in BRIP1 are associated with breast cancer risk: implications for gene panel testing. Journal of Medical Genetics, 2016, 53, 298-309.	1.5	94
30	Development and validation of a serum microRNA biomarker panel for detecting gastric cancer in a high-risk population. Gut, 2021, 70, 829-837.	6.1	94
31	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. Nature Communications, 2020, 11, 3833.	5.8	88
32	Common genetic determinants of breast-cancer risk in East Asian women: a collaborative study of 23 637 breast cancer cases and 25 579 controls. Human Molecular Genetics, 2013, 22, 2539-2550.	1.4	86
33	Health-related quality of life in Asian patients with breast cancer: a systematic review. BMJ Open, 2018, 8, e020512.	0.8	83
34	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast–ovarian cancer susceptibility locus. Nature Communications, 2016, 7, 12675.	5.8	78
35	Fine-Scale Mapping of the 5q11.2 Breast Cancer Locus Reveals at Least Three Independent Risk Variants Regulating MAP3K1. American Journal of Human Genetics, 2015, 96, 5-20.	2.6	76
36	Genetic implications of bilateral breast cancer: a population based cohort study. Lancet Oncology, The, 2005, 6, 377-382.	5.1	75

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37	HDAC1 and HDAC2 independently predict mortality in hepatocellular carcinoma by a competing risk regression model in a Southeast Asian population. Oncology Reports, 2015, 34, 2238-2250.	1.2	75
38	<i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. Cancer Research, 2017, 77, 2789-2799.	0.4	75
39	Positive impact of oral hydroxychloroquine and povidone-iodine throat spray for COVID-19 prophylaxis: An open-label randomized trial. International Journal of Infectious Diseases, 2021, 106, 314-322.	1.5	75
40	Familial concordance in cancer survival: a Swedish population-based study. Lancet Oncology, The, 2007, 8, 1001-1006.	5.1	69
41	Ethnic Differences in Survival after Breast Cancer in South East Asia. PLoS ONE, 2012, 7, e30995.	1.1	68
42	Splenectomy and the Risk of Sepsis. Annals of Surgery, 2014, 260, 1081-1087.	2.1	65
43	Prospective associations of appetitive traits at 3 and 12 months of age with body mass index and weight gain in the first 2 years of life. BMC Pediatrics, 2015, 15, 153.	0.7	60
44	Evidence that the 5p12 Variant rs10941679 Confers Susceptibility to Estrogen-Receptor-Positive Breast Cancer through FGF10 and MRPS30 Regulation. American Journal of Human Genetics, 2016, 99, 903-911.	2.6	59
45	Impact of breast surgery on survival in women presenting with metastatic breast cancer. British Journal of Surgery, 2011, 98, 1566-1572.	0.1	58
46	The incidence of breast cancer and changes in the use of hormone replacement therapy: A review of the evidence. Maturitas, 2009, 64, 80-85.	1.0	54
47	Is breast cancer prognosis inherited?. Breast Cancer Research, 2007, 9, R39.	2.2	53
48	Adjuvant! Online is overoptimistic in predicting survival of Asian breast cancer patients. European Journal of Cancer, 2012, 48, 982-989.	1.3	53
49	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. Human Molecular Genetics, 2014, 23, 6096-6111.	1.4	53
50	Prediction of breast cancer risk based on common genetic variants in women of East Asian ancestry. Breast Cancer Research, 2016, 18, 124.	2.2	52
51	Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. International Journal of Cancer, 2016, 139, 1303-1317.	2.3	51
52	Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes. JAMA Oncology, 2022, 8, e216744.	3.4	51
53	Is arterial base deficit still a useful prognostic marker in trauma? A systematic review. American Journal of Emergency Medicine, 2016, 34, 626-635.	0.7	50
54	Improving Outcomes in Breast Cancer for Low and Middle Income Countries. World Journal of Surgery, 2015, 39, 686-692.	0.8	47

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55	Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. Nature Communications, 2020, 11, 1217.	5.8	46
56	Impact of delayed treatment in women diagnosed with breast cancer: A population‐based study. Cancer Medicine, 2020, 9, 2435-2444.	1.3	46
57	Change of mammographic density predicts the risk of contralateral breast cancer - a case-control study. Breast Cancer Research, 2013, 15, R57.	2.2	44
58	The Predictive Accuracy of PREDICT. Medicine (United States), 2015, 94, e593.	0.4	42
59	Breast-conserving surgery*versus*mastectomy in young women with breast cancer in Asian settings. BJS Open, 2019, 3, 48-55.	0.7	42
60	Barriers to early presentation of self-discovered breast cancer in Singapore and Malaysia: a qualitative multicentre study. BMJ Open, 2015, 5, e009863.	0.8	41
61	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. Human Molecular Genetics, 2015, 24, 2966-2984.	1.4	40
62	Molecular epidemiology and its current clinical use in cancer management. Lancet Oncology, The, 2010, 11, 383-390.	5.1	39
63	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. American Journal of Human Genetics, 2020, 107, 837-848.	2.6	39
64	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. Human Molecular Genetics, 2015, 24, 285-298.	1.4	38
65	Evaluating genetic variants associated with breast cancer risk in high and moderate-penetrance genes in Asians. Carcinogenesis, 2017, 38, 511-518.	1.3	38
66	Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression. American Journal of Human Genetics, 2015, 97, 22-34.	2.6	37
67	The genetic interplay between body mass index, breast size and breast cancer risk: a Mendelian randomization analysis. International Journal of Epidemiology, 2019, 48, 781-794.	0.9	37
68	Birth rates among female cancer survivors. Cancer, 2013, 119, 1892-1899.	2.0	36
69	An intergenic risk locus containing an enhancer deletion in 2q35 modulates breast cancer risk by deregulating IGFBP5 expression. Human Molecular Genetics, 2016, 25, 3863-3876.	1.4	33
70	Significant Effect of Polymorphisms in*CYP2D6*on Response to Tamoxifen Therapy for Breast Cancer: A Prospective Multicenter Study. Clinical Cancer Research, 2017, 23, 2019-2026.	3.2	33
71	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. Breast Cancer Research, 2016, 18, 64.	2.2	31
72	Breast Cancer in South East Asia: Comparison of Presentation and Outcome Between a Middle Income and a High Income Country. World Journal of Surgery, 2012, 36, 2838-2846.	0.8	30

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73	Breast cancer risk assessment using genetic variants and risk factors in a Singapore Chinese population. Breast Cancer Research, 2014, 16, R64.	2.2	30
74	Parental perceptions of childhood seasonal influenza vaccination in Singapore: A cross-sectional survey. Vaccine, 2017, 35, 6096-6102.	1.7	29
75	Prognostic role of adjuvant radiotherapy in triple‐negative breast cancer: A historical cohort study. International Journal of Cancer, 2015, 137, 2504-2512.	2.3	28
76	Prevalence of PALB2 Mutations in Breast Cancer Patients in Multi-Ethnic Asian Population in Malaysia and Singapore. PLoS ONE, 2013, 8, e73638.	1.1	27
77	Polygenic risk scores for prediction of breast cancer risk in Asian populations. Genetics in Medicine, 2022, 24, 586-600.	1.1	27
78	Associations between workability and patient-reported physical, psychological and social outcomes in breast cancer survivors: a cross-sectional study. Supportive Care in Cancer, 2018, 26, 2815-2824.	1.0	25
79	High expression of intratumoral stromal proteins is associated with chemotherapy resistance in breast cancer. Oncotarget, 2016, 7, 55155-55168.	0.8	25
80	Trends in presentation, management and survival of patients with de novo metastatic breast cancer in a Southeast Asian setting. Scientific Reports, 2015, 5, 16252.	1.6	24
81	Fine-Scale Mapping of the 4q24 Locus Identifies Two Independent Loci Associated with Breast Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1680-1691.	1.1	24
82	Predicting Survival of De Novo Metastatic Breast Cancer in Asian Women: Systematic Review and Validation Study. PLoS ONE, 2014, 9, e93755.	1.1	24
83	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. European Journal of Human Genetics, 2022, 30, 349-362.	1.4	23
84	SILC for SILC: Single Institution Learning Curve for Single-Incision Laparoscopic Cholecystectomy. Minimally Invasive Surgery, 2013, 2013, 1-7.	0.1	22
85	Use of Activated Recombinant Factor VII in Severe Bleeding Â— Evidence for Efficacy and Safety in Trauma, Postpartum Hemorrhage, Cardiac Surgery, and Gastrointestinal Bleeding. Transfusion Medicine and Hemotherapy, 2012, 39, 139-150.	0.7	21
86	FGF receptor genes and breast cancer susceptibility: results from the Breast Cancer Association Consortium. British Journal of Cancer, 2014, 110, 1088-1100.	2.9	21
87	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. Cancer Causes and Control, 2016, 27, 679-693.	0.8	21
88	Development and validation of a circulating microRNA panel for the early detection of breast cancer. British Journal of Cancer, 2022, 126, 472-481.	2.9	21
89	Laparoscopic drainage of cryptogenic liver abscess. Surgical Endoscopy and Other Interventional Techniques, 2013, 27, 3308-3314.	1.3	20
90	Quantifying the natural history of breast cancer. British Journal of Cancer, 2013, 109, 2035-2043.	2.9	20

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91	Mammographic Breast Density and Common Genetic Variants in Breast Cancer Risk Prediction. PLoS ONE, 2015, 10, e0136650.	1.1	20
92	Differences in late cardiovascular mortality following acute myocardial infarction in three major Asian ethnic groups. European Heart Journal: Acute Cardiovascular Care, 2014, 3, 354-362.	0.4	19
93	Fine scale mapping of the 17q22 breast cancer locus using dense SNPs, genotyped within the Collaborative Oncological Gene-Environment Study (COGs). Scientific Reports, 2016, 6, 32512.	1.6	19
94	International Consortium on Mammographic Density: Methodology and population diversity captured across 22 countries. Cancer Epidemiology, 2016, 40, 141-151.	0.8	19
95	The <i>BRCA2</i> c.68-7T>A variant is not pathogenic: A model for clinical calibration of spliceogenicity. Human Mutation, 2018, 39, 729-741.	1.1	19
96	Determinants of breast size in Asian women. Scientific Reports, 2018, 8, 1201.	1.6	19
97	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 623-642.	1.1	19
98	High-Throughput Mutation Profiling Changes before and 3 Weeks after Chemotherapy in Newly Diagnosed Breast Cancer Patients. PLoS ONE, 2015, 10, e0142466.	1.1	19
99	Breast cancer risks associated with missense variants in breast cancer susceptibility genes. Genome Medicine, 2022, 14, 51.	3.6	19
100	Reliability of Ultrasound Duplex for Detection of Hemodynamically Significant Stenosis in Hemodialysis Access. Annals of Vascular Diseases, 2013, 6, 57-61.	0.2	18
101	Mammographic density assessed on paired raw and processed digital images and on paired screen-film and digital images across three mammography systems. Breast Cancer Research, 2016, 18, 130.	2.2	17
102	Effect of childbirth after treatment on long-term survival from breast cancer. British Journal of Surgery, 2010, 97, 1253-1259.	0.1	16
103	Estrogen Receptor Status in Relation to Risk of Contralateral Breast Cancer–A Population-Based Cohort Study. PLoS ONE, 2012, 7, e46535.	1.1	16
104	Demographic and Spatial Predictors of Anemia in Women of Reproductive Age in Timor-Leste: Implications for Health Program Prioritization. PLoS ONE, 2014, 9, e91252.	1.1	16
105	Validation of the CancerMath prognostic tool for breast cancer in Southeast Asia. BMC Cancer, 2016, 16, 820.	1.1	16
106	Outcome after neoadjuvant chemotherapy in Asian breast cancer patients. Cancer Medicine, 2017, 6, 173-185.	1.3	16
107	The association between weight at birth and breast cancer risk revisited using Mendelian randomisation. European Journal of Epidemiology, 2019, 34, 591-600.	2.5	16
108	Development of a microRNA Panel for Classification of Abnormal Mammograms for Breast Cancer. Cancers, 2021, 13, 2130.	1.7	16

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109	Phase Ib/II randomized, open-label study of doxorubicin and cyclophosphamide with or without low-dose, short-course sunitinib in the pre-operative treatment of breast cancer. Oncotarget, 2016, 7, 64089-64099.	0.8	16
110	Breast cancer prognosis is inherited independently of patient, tumor and treatment characteristics. International Journal of Cancer, 2012, 130, 2103-2110.	2.3	15
111	Performance of Single-Nucleotide Polymorphisms in Breast Cancer Risk Prediction Models: A Systematic Review and Meta-analysis. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 506-521.	1.1	15
112	Does the Axillary Lymph Node Ratio Have Any Added Prognostic Value over pN Staging for South East Asian Breast Cancer Patients?. PLoS ONE, 2012, 7, e45809.	1.1	14
113	Prognostic implications of estrogen receptor pattern of both tumors in contralateral breast cancer. Breast Cancer Research and Treatment, 2012, 134, 793-800.	1.1	14
114	Genetic variation at CYP3A is associated with age at menarche and breast cancer risk: a case-control study. Breast Cancer Research, 2014, 16, R51.	2.2	14
115	Birth rates among male cancer survivors and mortality rates among their offspring: a population-based study from Sweden. BMC Cancer, 2016, 16, 196.	1.1	14
116	Re-evaluating genetic variants identified in candidate gene studies of breast cancer risk using data from nearly 280,000 women of Asian and European ancestry. EBioMedicine, 2019, 48, 203-211.	2.7	14
117	Incidence of breast cancer attributable to breast density, modifiable and non-modifiable breast cancer risk factorsÂin Singapore. Scientific Reports, 2020, 10, 503.	1.6	14
118	Impact of deviation from guideline recommended treatment on breast cancer survival in Asia. Scientific Reports, 2020, 10, 1330.	1.6	14
119	Predicting the Likelihood of Carrying a <i>BRCA1</i> or <i>BRCA2</i> Mutation in Asian Patients With Breast Cancer. Journal of Clinical Oncology, 2022, 40, 1542-1551.	0.8	14
120	Prognostic value of axillary lymph node status after neoadjuvant chemotherapy. Results from a multicentre study. European Journal of Cancer, 2011, 47, 1186-1192.	1.3	13
121	Singapore's anatomical future: Quo Vadis?. Anatomical Sciences Education, 2012, 5, 234-240.	2.5	13
122	Inter-Ethnic Differences in Quantified Coronary Artery Disease Severity and All-Cause Mortality among Dutch and Singaporean Percutaneous Coronary Intervention Patients. PLoS ONE, 2015, 10, e0131977.	1.1	13
123	Women&rsquo;s preferences, willingness-to-pay, and predicted uptake for single-nucleotide polymorphism gene testing to guide personalized breast cancer screening strategies: a discrete choice experiment. Patient Preference and Adherence, 2018, Volume 12, 1837-1852.	0.8	13
124	Comparison of self-reported and register-based hospital medical data on comorbidities in women. Scientific Reports, 2019, 9, 3527.	1.6	13
125	Presentation of breast cancer, help seeking behaviour and experience of patients in their cancer journey in Singapore: a qualitative study. BMC Cancer, 2020, 20, 1080.	1.1	13
126	Ability to predict breast cancer in Asian women using a polygenic susceptibility model. Breast Cancer Research and Treatment, 2011, 127, 805-812.	1.1	12

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127	Large-scale genotyping identifies a new locus at 22q13.2 associated with female breast size. Journal of Medical Genetics, 2013, 50, 666-673.	1.5	12
128	Spectrum of very early breast cancer in a setting without organised screening. British Journal of Cancer, 2014, 110, 2187-2194.	2.9	12
129	Assessment of a fully automated, high-throughput mammographic density measurement tool for use with processed digital mammograms. Cancer Causes and Control, 2014, 25, 1037-1043.	0.8	12
130	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. PLoS ONE, 2016, 11, e0160316.	1.1	12
131	Restructuring the surgical service during the COVID-19 pandemic: Experience from a tertiary institution in Singapore. American Journal of Surgery, 2020, 220, 553-555.	0.9	12
132	Breast Cancer Onset in Twins and Women With Bilateral Disease. Journal of Clinical Oncology, 2008, 26, 4086-4091.	0.8	11
133	Recurrent mutation testing of BRCA1 and BRCA2 in Asian breast cancer patients identify carriers in those with presumed low risk by family history. Breast Cancer Research and Treatment, 2014, 144, 635-642.	1.1	11
134	Cost effectiveness analysis of a polygenic risk tailored breast cancer screening programme in Singapore. BMC Health Services Research, 2021, 21, 379.	0.9	11
135	Cohort profile: The Singapore Breast Cancer Cohort (SGBCC), a multi-center breast cancer cohort for evaluation of phenotypic risk factors and genetic markers. PLoS ONE, 2021, 16, e0250102.	1.1	11
136	BREAst screening Tailored for HEr (BREATHE)â€"A study protocol on personalised risk-based breast cancer screening programme. PLoS ONE, 2022, 17, e0265965.	1.1	11
137	Interest and attitudes of patients, cancer physicians, medical students and cancer researchers towards a spectrum of genetic tests relevant to breast cancer patients. Breast, 2013, 22, 47-52.	0.9	10
138	The impact of in situ breast cancer and family history on risk of subsequent breast cancer events and mortality - a population-based study from Sweden. Breast Cancer Research, 2016, 18, 105.	2.2	10
139	Zinc and vitamin C intake increases spike and neutralising antibody production following SARSâ€CoVâ€2 infection. Clinical and Translational Medicine, 2022, 12, e731.	1.7	10
140	The functional ALDH2 polymorphism is associated with breast cancer risk: A pooled analysis from the Breast Cancer Association Consortium. Molecular Genetics & Genomic Medicine, 2019, 7, e707.	0.6	9
141	Factors associated with false-positive mammography at first screen in an Asian population. PLoS ONE, 2019, 14, e0213615.	1.1	9
142	Cost-effectiveness Analysis of Breast Cancer Screening Using Mammography in Singapore: A Modeling Study. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 653-660.	1.1	9
143	Overlap of high-risk individuals predicted by family history, and genetic and non-genetic breast cancer risk prediction models: implications for risk stratification. BMC Medicine, 2022, 20, 150.	2.3	9
144	Mortality among offspring of women diagnosed with cancer: A populationâ€based cohort study. International Journal of Cancer, 2013, 132, 2432-2438.	2.3	8

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145	Determinants of satisfaction with cosmetic outcome in breast cancer survivors: A cross-sectional study. PLoS ONE, 2018, 13, e0193099.	1.1	8
146	Immunohistochemistry study of tumor vascular normalization and anti-angiogenic effects of sunitinib versus bevacizumab prior to dose-dense doxorubicin/cyclophosphamide chemotherapy in HER2-negative breast cancer. Breast Cancer Research and Treatment, 2022, 192, 131-142.	1.1	8
147	DNA methylation and breast cancer-associated variants. Breast Cancer Research and Treatment, 2021, 188, 713-727.	1.1	7
148	Is Genetic Background Important in Lung Cancer Survival?. PLoS ONE, 2009, 4, e5588.	1.1	6
149	A comparative population-based study of prostate cancer incidence and mortality rates in Singapore, Sweden and Geneva, Switzerland from 1973 to 2006. BMC Cancer, 2012, 12, 222.	1.1	6
150	The impact of breast cancer-specific birth cohort effects among younger and older Chinese populations. International Journal of Cancer, 2016, 139, 527-534.	2.3	6
151	Ultrasound Guided Optoacoustic Tomography in Assessment of Tumor Margins for Lumpectomies. Translational Oncology, 2020, 13, 254-261.	1.7	6
152	Pleiotropy-guided transcriptome imputation from normal and tumor tissues identifies candidate susceptibility genes for breast and ovarian cancer. Human Genetics and Genomics Advances, 2021, 2, 100042.	1.0	6
153	Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. American Journal of Human Genetics, 2021, 108, 1190-1203.	2.6	6
154	Breast cancer risk stratification for mammographic screening: A nation–wide screening cohort of 24,431 women in Singapore. Cancer Medicine, 2021, 10, 8182-8191.	1.3	6
155	The association of age at menarche and adult height with mammographic density in the International Consortium of Mammographic Density. Breast Cancer Research, 2022, 24, .	2.2	6
156	Prognostic information of a previously diagnosed sister is an independent prognosticator for a newly diagnosed sister with breast cancer. Annals of Oncology, 2014, 25, 1966-1972.	0.6	5
157	Two truncating variants in FANCC and breast cancer risk. Scientific Reports, 2019, 9, 12524.	1.6	5
158	Epidemiological and ES cell–based functional evaluation of BRCA2 variants identified in families with breast cancer. Human Mutation, 2021, 42, 200-212.	1.1	4
159	Characterisation of protein-truncating and missense variants in PALB2 in 15 768 women from Malaysia and Singapore. Journal of Medical Genetics, 2021, , jmedgenet-2020-107471.	1.5	4
160	Gene-Environment Interactions Relevant to Estrogen and Risk of Breast Cancer: Can Gene-Environment Interactions Be Detected Only among Candidate SNPs from Genome-Wide Association Studies?. Cancers, 2021, 13, 2370.	1.7	4
161	Impact of older age on presentation, management and outcome of breast cancer in the multi-ethnic Asian population of Singapore. Journal of Geriatric Oncology, 2011, 2, 50-57.	0.5	3
162	Prediction of Lymph Node Involvement in Patients with Breast Tumors Measuring 3–5 cm in a Middle–Income Setting: the Role of CancerMath. World Journal of Surgery, 2014, 38, 3133-3137.	0.8	3

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163	Using peer review to distribute group work marks equitably between medical students. BMC Medical Education, 2017, 17, 172.	1.0	3
164	Aortic Wall Extracellular Matrix Proteins Correlate with Syntax Score in Patients Undergoing Coronary Artery Bypass Surgery. Open Cardiovascular Medicine Journal, 2016, 10, 48-56.	0.6	3
165	Germline breast cancer susceptibility genes, tumor characteristics, and survival. Genome Medicine, 2021, 13, 185.	3.6	3
166	A constant risk for familial breast cancer? A population-based family study. Breast Cancer Research, 2009, 11, R30.	2.2	2
167	rs2735383, located at a microRNA binding site in the 3′UTR of NBS1, is not associated with breast cancer risk. Scientific Reports, 2016, 6, 36874.	1.6	2
168	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. Scientific Reports, 2020, 10, 9688.	1.6	2
169	Phase II study of neoadjuvant weekly paclitaxel and carboplatin with lapatinib in HER2+ breast cancer.. Journal of Clinical Oncology, 2014, 32, 619-619.	0.8	2
170	Association between Breast Cancer Polygenic Risk Score and Chemotherapy-Induced Febrile Neutropenia: Null Results. Cancers, 2022, 14, 2714.	1.7	2
171	Diagnostic work-up of contralateral breast cancers has not improved over calendar period. Breast Cancer Research and Treatment, 2010, 122, 889-895.	1.1	1
172	Influence of radiotherapy for the first tumor on aggressiveness of contralateral breast cancer. International Journal of Cancer, 2013, 132, 2388-2394.	2.3	1
173	Do thrombotic events during endovascular interventions lead to poorer outcomes in patients with severe limb ischemia?. Vascular, 2015, 23, 245-252.	0.4	1
174	A genome-wide association study identifies three novel genetic markers for response to tamoxifen: A prospective multicenter study. PLoS ONE, 2018, 13, e0201606.	1.1	1
175	Abstract 2780: Evaluation of genetic variants in high and moderate-penetrance breast cancer susceptibility genes in East Asians. , 2015, , .		1
176	Abstract 2031: Association between CYP2D6 genotype and response to tamoxifen in a prospective multicenter study in Japan. , 2016, , .		1
177	Genomic predictors of neoadjuvant chemotherapy (NACT) response in breast cancer (BC).. Journal of Clinical Oncology, 2017, 35, e12122-e12122.	0.8	1
178	Associations between Pre-Diagnostic Physical Activity with Breast Cancer Characteristics and Survival. Cancers, 2022, 14, 1756.	1.7	1
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