

Jingmei Li

List of Publications by Year in descending order

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Version: 2024-02-01

172
papers

12,182
citations

44069

48
h-index

32842

100
g-index

181
all docs

181
docs citations

181
times ranked

17329
citing authors

#	ARTICLE	IF	CITATIONS
1	Interval breast cancer is associated with interferon immune response. <i>European Journal of Cancer</i> , 2022, 162, 194-205.	2.8	3
2	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 349-362.	2.8	23
3	Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes. <i>JAMA Oncology</i> , 2022, 8, e216744.	7.1	51
4	Predicting the Likelihood of Carrying a <i>BRCA1</i> or <i>BRCA2</i> Mutation in Asian Patients With Breast Cancer. <i>Journal of Clinical Oncology</i> , 2022, 40, 1542-1551.	1.6	14
5	BREast screening Tailored for HER (BREATHE) – A study protocol on personalised risk-based breast cancer screening programme. <i>PLoS ONE</i> , 2022, 17, e0265965.	2.5	11
6	Associations between Pre-Diagnostic Physical Activity with Breast Cancer Characteristics and Survival. <i>Cancers</i> , 2022, 14, 1756.	3.7	1
7	Polygenic risk scores for prediction of breast cancer risk in Asian populations. <i>Genetics in Medicine</i> , 2022, 24, 586-600.	2.4	27
8	Dual-RNA controlled delivery system inhibited tumor growth by apoptosis induction and TME activation. <i>Journal of Controlled Release</i> , 2022, 344, 97-112.	9.9	4
9	Overlap of high-risk individuals predicted by family history, and genetic and non-genetic breast cancer risk prediction models: implications for risk stratification. <i>BMC Medicine</i> , 2022, 20, 150.	5.5	9
10	Relevance of the MHC region for breast cancer susceptibility in Asians. <i>Breast Cancer</i> , 2022, 29, 869-879.	2.9	1
11	Breast cancer risks associated with missense variants in breast cancer susceptibility genes. <i>Genome Medicine</i> , 2022, 14, 51.	8.2	19
12	Association between Breast Cancer Polygenic Risk Score and Chemotherapy-Induced Febrile Neutropenia: Null Results. <i>Cancers</i> , 2022, 14, 2714.	3.7	2
13	Epidemiological and ES cell-based functional evaluation of BRCA2 variants identified in families with breast cancer. <i>Human Mutation</i> , 2021, 42, 200-212.	2.5	4
14	Association between breast cancer risk and disease aggressiveness: Characterizing underlying gene expression patterns. <i>International Journal of Cancer</i> , 2021, 148, 884-894.	5.1	3
15	Breast Cancer Risk Genes – Association Analysis in More than 113,000 Women. <i>New England Journal of Medicine</i> , 2021, 384, 428-439.	27.0	532
16	DNA methylation and breast cancer-associated variants. <i>Breast Cancer Research and Treatment</i> , 2021, 188, 713-727.	2.5	7
17	Characterisation of protein-truncating and missense variants in PALB2 in 15 768 women from Malaysia and Singapore. <i>Journal of Medical Genetics</i> , 2021, , jmedgenet-2020-107471.	3.2	4
18	Cohort profile: The Singapore Breast Cancer Cohort (SGBCC), a multi-center breast cancer cohort for evaluation of phenotypic risk factors and genetic markers. <i>PLoS ONE</i> , 2021, 16, e0250102.	2.5	11

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19	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?. <i>Cancers</i> , 2021, 13, 2370.	3.7	4
20	Pleiotropy-guided transcriptome imputation from normal and tumor tissues identifies candidate susceptibility genes for breast and ovarian cancer. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100042.	1.7	6
21	Functionalized DMP-039 Hybrid Nanoparticle as a Novel mRNA Vector for Efficient Cancer Suicide Gene Therapy. <i>International Journal of Nanomedicine</i> , 2021, Volume 16, 5211-5232.	6.7	24
22	Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. <i>American Journal of Human Genetics</i> , 2021, 108, 1190-1203.	6.2	6
23	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	27.8	183
24	Single Micelle Vectors based on Lipid/Block Copolymer Compositions as mRNA Formulations for Efficient Cancer Immunogene Therapy. <i>Molecular Pharmaceutics</i> , 2021, 18, 4029-4045.	4.6	13
25	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 623-642.	2.5	19
26	Breast cancer risk stratification for mammographic screening: A nationwide screening cohort of 24,431 women in Singapore. <i>Cancer Medicine</i> , 2021, 10, 8182-8191.	2.8	6
27	Germline breast cancer susceptibility genes, tumor characteristics, and survival. <i>Genome Medicine</i> , 2021, 13, 185.	8.2	3
28	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. <i>Nature Communications</i> , 2020, 11, 3833.	12.8	88
29	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.	6.2	39
30	<p></p>Treatment of Colon Cancer by Degradable rrPPC Nano-Conjugates Delivered STAT3 siRNA</p>. <i>International Journal of Nanomedicine</i> , 2020, Volume 15, 9875-9890.	6.7	22
31	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , 2020, 10, 9688.	3.3	2
32	Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. <i>Nature Communications</i> , 2020, 11, 1217.	12.8	46
33	Impact of delayed treatment in women diagnosed with breast cancer: A population-based study. <i>Cancer Medicine</i> , 2020, 9, 2435-2444.	2.8	46
34	Incidence of breast cancer attributable to breast density, modifiable and non-modifiable breast cancer risk factors in Singapore. <i>Scientific Reports</i> , 2020, 10, 503.	3.3	14
35	Impact of deviation from guideline recommended treatment on breast cancer survival in Asia. <i>Scientific Reports</i> , 2020, 10, 1330.	3.3	14
36	The genetic interplay between body mass index, breast size and breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2019, 48, 781-794.	1.9	37

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37	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. <i>EBioMedicine</i> , 2019, 48, 203-211.	6.1	14
38	Two truncating variants in <i>FANCC</i> and breast cancer risk. <i>Scientific Reports</i> , 2019, 9, 12524.	3.3	5
39	Joint association of mammographic density adjusted for age and body mass index and polygenic risk score with breast cancer risk. <i>Breast Cancer Research</i> , 2019, 21, 68.	5.0	31
40	The long-term prognostic and predictive capacity of cyclin D1 gene amplification in 2305 breast tumours. <i>Breast Cancer Research</i> , 2019, 21, 34.	5.0	48
41	Comparison of self-reported and register-based hospital medical data on comorbidities in women. <i>Scientific Reports</i> , 2019, 9, 3527.	3.3	13
42	Factors associated with false-positive mammography at first screen in an Asian population. <i>PLoS ONE</i> , 2019, 14, e0213615.	2.5	9
43	The association between weight at birth and breast cancer risk revisited using Mendelian randomisation. <i>European Journal of Epidemiology</i> , 2019, 34, 591-600.	5.7	16
44	Prevalence of <i>BRCA1</i> and <i>BRCA2</i> pathogenic variants in a large, unselected breast cancer cohort. <i>International Journal of Cancer</i> , 2019, 144, 1195-1204.	5.1	31
45	A comprehensive tool for measuring mammographic density changes over time. <i>Breast Cancer Research and Treatment</i> , 2018, 169, 371-379.	2.5	45
46	Determinants of breast size in Asian women. <i>Scientific Reports</i> , 2018, 8, 1201.	3.3	19
47	Family History, Reproductive, and Lifestyle Risk Factors for Fibroadenoma and Breast Cancer. <i>JNCI Cancer Spectrum</i> , 2018, 2, pky051.	2.9	19
48	Differential Burden of Rare and Common Variants on Tumor Characteristics, Survival, and Mode of Detection in Breast Cancer. <i>Cancer Research</i> , 2018, 78, 6329-6338.	0.9	19
49	Inherited factors contribute to an inverse association between preeclampsia and breast cancer. <i>Breast Cancer Research</i> , 2018, 20, 6.	5.0	14
50	Common genetic variation and novel loci associated with volumetric mammographic density. <i>Breast Cancer Research</i> , 2018, 20, 30.	5.0	18
51	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , 2018, 50, 968-978.	21.4	184
52	<i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017, 77, 2789-2799.	0.9	75
53	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. <i>Nature Genetics</i> , 2017, 49, 834-841.	21.4	426
54	Assessment of Breast Cancer Risk Factors Reveals Subtype Heterogeneity. <i>Cancer Research</i> , 2017, 77, 3708-3717.	0.9	87

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55	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	27.8	1,099
56	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	21.4	289
57	Common shared genetic variation behind decreased risk of breast cancer in celiac disease. <i>Scientific Reports</i> , 2017, 7, 5942.	3.3	5
58	Longitudinal fluctuation in mammographic percent density differentiates between interval and screen-detected breast cancer. <i>International Journal of Cancer</i> , 2017, 140, 34-40.	5.1	6
59	Molecular Differences between Screen-Detected and Interval Breast Cancers Are Largely Explained by PAM50 Subtypes. <i>Clinical Cancer Research</i> , 2017, 23, 2584-2592.	7.0	15
60	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017, 19, 599-603.	2.4	67
61	Associations between childhood body size and seventeen adverse outcomes: analysis of 65,057 European women. <i>Scientific Reports</i> , 2017, 7, 16917.	3.3	8
62	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017, 46, 1814-1822.	1.9	45
63	Risk and predictors of psoriasis in patients with breast cancer: a Swedish population-based cohort study. <i>BMC Medicine</i> , 2017, 15, 154.	5.5	13
64	Body size in early life and risk of breast cancer. <i>Breast Cancer Research</i> , 2017, 19, 84.	5.0	15
65	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. <i>Breast Cancer Research</i> , 2017, 19, 119.	5.0	43
66	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. <i>PLoS Medicine</i> , 2016, 13, e1002105.	8.4	118
67	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. <i>PLoS ONE</i> , 2016, 11, e0160316.	2.5	12
68	Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. <i>International Journal of Cancer</i> , 2016, 139, 1303-1317.	5.1	51
69	<i>PALB2</i> , <i>CHEK2</i> and <i>ATM</i> rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016, 53, 800-811.	3.2	174
70	Patient survival and tumor characteristics associated with CHEK2:p.I157T â€” findings from the Breast Cancer Association Consortium. <i>Breast Cancer Research</i> , 2016, 18, 98.	5.0	39
71	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. <i>Breast Cancer Research</i> , 2016, 18, 64.	5.0	31
72	Genetic predisposition to ductal carcinoma in situ of the breast. <i>Breast Cancer Research</i> , 2016, 18, 22.	5.0	43

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73	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. <i>Cancer Causes and Control</i> , 2016, 27, 679-693.	1.8	21
74	Five endometrial cancer risk loci identified through genome-wide association analysis. <i>Nature Genetics</i> , 2016, 48, 667-674.	21.4	77
75	Evidence that the 5p12 Variant rs10941679 Confers Susceptibility to Estrogen-Receptor-Positive Breast Cancer through FGF10 and MRPS30 Regulation. <i>American Journal of Human Genetics</i> , 2016, 99, 903-911.	6.2	59
76	Worse quality of life in young and recently diagnosed breast cancer survivors compared with female survivors of other cancers: A cross-sectional study. <i>International Journal of Cancer</i> , 2016, 139, 2415-2425.	5.1	23
77	Genetic Risk Score Mendelian Randomization Shows that Obesity Measured as Body Mass Index, but not Waist:Hip Ratio, Is Causal for Endometrial Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1503-1510.	2.5	64
78	rs2735383, located at a microRNA binding site in the 3'UTR of NBS1, is not associated with breast cancer risk. <i>Scientific Reports</i> , 2016, 6, 36874.	3.3	2
79	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	12.8	78
80	Common diseases as determinants of menopausal age. <i>Human Reproduction</i> , 2016, 31, 2856-2864.	0.9	42
81	Assessing within-woman changes in mammographic density: a comparison of fully versus semi-automated area-based approaches. <i>Cancer Causes and Control</i> , 2016, 27, 481-491.	1.8	15
82	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2016, 23, 77-91.	3.1	62
83	Variants in 6q25.1 Are Associated with Mammographic Density in Malaysian Chinese Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 327-333.	2.5	10
84	Genetic variation in the immunosuppression pathway genes and breast cancer susceptibility: a pooled analysis of 42,510 cases and 40,577 controls from the Breast Cancer Association Consortium. <i>Human Genetics</i> , 2016, 135, 137-154.	3.8	8
85	Associations of Breast Cancer Risk Prediction Tools With Tumor Characteristics and Metastasis. <i>Journal of Clinical Oncology</i> , 2016, 34, 251-258.	1.6	45
86	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. <i>Journal of the National Cancer Institute</i> , 2016, 108, djv315.	6.3	77
87	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	1.4	18
88	RAD51B in Familial Breast Cancer. <i>PLoS ONE</i> , 2016, 11, e0153788.	2.5	26
89	A Combined Segmentation and Registration Framework for Bilateral and Temporal Mammogram Analysis. <i>Journal of Medical Imaging and Health Informatics</i> , 2016, 6, 380-386.	0.3	2
90	Investigation of gene-environment interactions between 47 newly identified breast cancer susceptibility loci and environmental risk factors. <i>International Journal of Cancer</i> , 2015, 136, E685-96.	5.1	34

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91	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. Scientific Reports, 2015, 5, 17369.	3.3	35
92	A polymorphism in the base excision repair gene PARP2 is associated with differential prognosis by chemotherapy among postmenopausal breast cancer patients. BMC Cancer, 2015, 15, 978.	2.6	6
93	Common germline polymorphisms associated with breast cancer-specific survival. Breast Cancer Research, 2015, 17, 58.	5.0	26
94	Identification of two novel mammographic density loci at 6Q25.1. Breast Cancer Research, 2015, 17, 75.	5.0	24
95	A comprehensive evaluation of interaction between genetic variants and use of menopausal hormone therapy on mammographic density. Breast Cancer Research, 2015, 17, 110.	5.0	19
96	Large-Scale Genomic Analyses Link Reproductive Aging to Hypothalamic Signaling, Breast Cancer Susceptibility, and BRCA1-Mediated DNA Repair. Obstetrical and Gynecological Survey, 2015, 70, 758-762.	0.4	0
97	The Genetic Basis of Quality of Life in Healthy Swedish Women: A Candidate Gene Approach. PLoS ONE, 2015, 10, e0118292.	2.5	8
98	Mammographic Breast Density and Common Genetic Variants in Breast Cancer Risk Prediction. PLoS ONE, 2015, 10, e0136650.	2.5	20
99	SNP-SNP interaction analysis of NF- κ B signaling pathway on breast cancer survival. Oncotarget, 2015, 6, 37979-37994.	1.8	20
100	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. Journal of the National Cancer Institute, 2015, 107, .	6.3	428
101	The association between lower educational attainment and depression owing to shared genetic effects? Results in ~25â€‰%000 subjects. Molecular Psychiatry, 2015, 20, 735-743.	7.9	59
102	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. Human Molecular Genetics, 2015, 24, 2966-2984.	2.9	40
103	Risk Factors and Tumor Characteristics of Interval Cancers by Mammographic Density. Journal of Clinical Oncology, 2015, 33, 1030-1037.	1.6	99
104	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.	6.2	76
105	Breast cancer genetic risk profile is differentially associated with interval and screen-detected breast cancers. Annals of Oncology, 2015, 26, 517-522.	1.2	38
106	Inherited variants in the inner centromere protein (INCENP) gene of the chromosomal passenger complex contribute to the susceptibility of ER-negative breast cancer. Carcinogenesis, 2015, 36, 256-271.	2.8	14
107	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. Nature Genetics, 2015, 47, 373-380.	21.4	513
108	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.	6.2	37

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109	Identification of Novel Genetic Markers of Breast Cancer Survival. Journal of the National Cancer Institute, 2015, 107, .	6.3	56
110	Candidate locus analysis of the TERT-CLPTM1L cancer risk region on chromosome 5p15 identifies multiple independent variants associated with endometrial cancer risk. Human Genetics, 2015, 134, 231-245.	3.8	34
111	Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with prognosis of estrogen receptor-negative breast cancer after chemotherapy. Breast Cancer Research, 2015, 17, 18.	5.0	20
112	Novel Associations between Common Breast Cancer Susceptibility Variants and Risk-Predicting Mammographic Density Measures. Cancer Research, 2015, 75, 2457-2467.	0.9	55
113	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. Nature Genetics, 2015, 47, 1294-1303.	21.4	357
114	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. Journal of the National Cancer Institute, 2015, 107, djv219.	6.3	99
115	Comprehensive genetic assessment of the ESR1 locus identifies a risk region for endometrial cancer. Endocrine-Related Cancer, 2015, 22, 851-861.	3.1	25
116	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.	2.5	24
117	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.	2.9	38
118	Fine-mapping of the HNF1B multicancer locus identifies candidate variants that mediate endometrial cancer risk. Human Molecular Genetics, 2015, 24, 1478-1492.	2.9	50
119	Ethnic Differences in Mammographic Densities: An Asian Cross-Sectional Study. PLoS ONE, 2015, 10, e0117568.	2.5	44
120	In Search for the Genetic Basis of Quality of Life in Healthy Swedish Women- A GWAS Study Using the iCOGS Custom Genotyping Array. PLoS ONE, 2015, 10, e0140563.	2.5	2
121	MicroRNA Related Polymorphisms and Breast Cancer Risk. PLoS ONE, 2014, 9, e109973.	2.5	49
122	Area and Volumetric Density Estimation in Processed Full-Field Digital Mammograms for Risk Assessment of Breast Cancer. PLoS ONE, 2014, 9, e110690.	2.5	24
123	Automated Measurement of Volumetric Mammographic Density: A Tool for Widespread Breast Cancer Risk Assessment. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 1764-1772.	2.5	62
124	Digital mammographic density and breast cancer risk: a case-control study of six alternative density assessment methods. Breast Cancer Research, 2014, 16, 439.	5.0	165
125	Volumetric Mammographic Density: Heritability and Association With Breast Cancer Susceptibility Loci. Journal of the National Cancer Institute, 2014, 106, dju334-dju334.	6.3	21
126	Genetic Predisposition to In Situ and Invasive Lobular Carcinoma of the Breast. PLoS Genetics, 2014, 10, e1004285.	3.5	39

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127	Enhancement of Mammographic Density Measures in Breast Cancer Risk Prediction. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1314-1323.	2.5	17
128	2q36.3 is associated with prognosis for oestrogen receptor-negative breast cancer patients treated with chemotherapy. <i>Nature Communications</i> , 2014, 5, 4051.	12.8	16
129	Polygenic scores associated with educational attainment in adults predict educational achievement and ADHD symptoms in children. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2014, 165, 510-520.	1.7	40
130	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2014, 23, 6096-6111.	2.9	53
131	Prognostic information of a previously diagnosed sister is an independent prognosticator for a newly diagnosed sister with breast cancer. <i>Annals of Oncology</i> , 2014, 25, 1966-1972.	1.2	5
132	Automated breast tissue density assessment using high order regional texture descriptors in mammography. <i>Proceedings of SPIE</i> , 2014, , .	0.8	2
133	CYP2B6*6 is associated with increased breast cancer risk. <i>International Journal of Cancer</i> , 2014, 134, 426-430.	5.1	24
134	A large-scale assessment of two-way SNP interactions in breast cancer susceptibility using 46 450 cases and 42 461 controls from the breast cancer association consortium. <i>Human Molecular Genetics</i> , 2014, 23, 1934-1946.	2.9	32
135	Identification of New Genetic Susceptibility Loci for Breast Cancer Through Consideration of Gene-Environment Interactions. <i>Genetic Epidemiology</i> , 2014, 38, 84-93.	1.3	28
136	Genome-wide association study identifies multiple loci associated with both mammographic density and breast cancer risk. <i>Nature Communications</i> , 2014, 5, 5303.	12.8	109
137	Mammographic Density Phenotypes and Risk of Breast Cancer: A Meta-analysis. <i>Journal of the National Cancer Institute</i> , 2014, 106, .	6.3	261
138	FGF receptor genes and breast cancer susceptibility: results from the Breast Cancer Association Consortium. <i>British Journal of Cancer</i> , 2014, 110, 1088-1100.	6.4	21
139	Assessment of a fully automated, high-throughput mammographic density measurement tool for use with processed digital mammograms. <i>Cancer Causes and Control</i> , 2014, 25, 1037-1043.	1.8	12
140	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	12.8	105
141	Genetic variation in mitotic regulatory pathway genes is associated with breast tumor grade. <i>Human Molecular Genetics</i> , 2014, 23, 6034-6046.	2.9	12
142	The aetiology of convulsive status epilepticus: A study of 258 cases in Western China. <i>Seizure: the Journal of the British Epilepsy Association</i> , 2014, 23, 717-721.	2.0	12
143	Abstract 3275: Genome-wide association study of childhood body fatness as a risk factor of breast cancer. , 2014, , .		0
144	Fine-Scale Mapping of the FGFR2 Breast Cancer Risk Locus: Putative Functional Variants Differentially Bind FOXA1 and E2F1. <i>American Journal of Human Genetics</i> , 2013, 93, 1046-1060.	6.2	98

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145	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384.	21.4	493
146	Change of mammographic density predicts the risk of contralateral breast cancer - a case-control study. <i>Breast Cancer Research</i> , 2013, 15, R57.	5.0	44
147	Confirmation of the reduction of hormone replacement therapy-related breast cancer risk for carriers of the HSD17B1_937_G variant. <i>Breast Cancer Research and Treatment</i> , 2013, 138, 543-548.	2.5	10
148	Functional Variants at the 11q13 Risk Locus for Breast Cancer Regulate Cyclin D1 Expression through Long-Range Enhancers. <i>American Journal of Human Genetics</i> , 2013, 92, 489-503.	6.2	201
149	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	21.4	960
150	GWAS of 126,559 Individuals Identifies Genetic Variants Associated with Educational Attainment. <i>Science</i> , 2013, 340, 1467-1471.	12.6	750
151	A genome-wide association study to identify genetic susceptibility loci that modify ductal and lobular postmenopausal breast cancer risk associated with menopausal hormone therapy use: a two-stage design with replication. <i>Breast Cancer Research and Treatment</i> , 2013, 138, 529-542.	2.5	18
152	Mammographic Density Reduction Is a Prognostic Marker of Response to Adjuvant Tamoxifen Therapy in Postmenopausal Patients With Breast Cancer. <i>Journal of Clinical Oncology</i> , 2013, 31, 2249-2256.	1.6	113
153	Large-scale genotyping identifies a new locus at 22q13.2 associated with female breast size. <i>Journal of Medical Genetics</i> , 2013, 50, 666-673.	3.2	12
154	Calcium intake is not related to breast cancer risk among Singapore Chinese women. <i>International Journal of Cancer</i> , 2013, 133, 680-686.	5.1	15
155	Association of CYP2D6 metabolizer status with mammographic density change in response to tamoxifen treatment. <i>Breast Cancer Research</i> , 2013, 15, R93.	5.0	10
156	The UGT1A6_19_CG genotype is a breast cancer risk factor. <i>Frontiers in Genetics</i> , 2013, 4, 104.	2.3	8
157	Common Breast Cancer Susceptibility Variants in <i>LSP1</i> and <i>RAD51L1</i> Are Associated with Mammographic Density Measures that Predict Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1156-1166.	2.5	101
158	Identification of a novel percent mammographic density locus at 12q24. <i>Human Molecular Genetics</i> , 2012, 21, 3299-3305.	2.9	31
159	Mammographic Breast Density and Breast Cancer: Evidence of a Shared Genetic Basis. <i>Cancer Research</i> , 2012, 72, 1478-1484.	0.9	54
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