Jingmei Li

List of Publications by Year
in descending order

172 papers	12,182 citations	50566 48 h-index	37326 100 g-index
181 all docs	181 docs citations	181 times ranked	18966 citing authors

#	Article	IF	Citations
1	Interval breast cancer is associated with interferon immune response. European Journal of Cancer, 2022, 162, 194-205.	1.3	3
2	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. European Journal of Human Genetics, 2022, 30, 349-362.	1.4	23
3	Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes. JAMA Oncology, 2022, 8, e216744.	3.4	51
4	Predicting the Likelihood of Carrying a BRCA1 or BRCA2 Mutation in Asian Patients With Breast Cancer. Journal of Clinical Oncology, 2022, 40, 1542-1551.	0.8	14
5	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
6	Associations between Pre-Diagnostic Physical Activity with Breast Cancer Characteristics and Survival. Cancers, 2022, 14, 1756.	1.7	1
7	Polygenic risk scores for prediction of breast cancer risk in Asian populations. Genetics in Medicine, 2022, 24, 586-600.	1.1	27
8	Dual-RNA controlled delivery system inhibited tumor growth by apoptosis induction and TME activation. Journal of Controlled Release, 2022, 344, 97-112.	4.8	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. BMC Medicine, 2022, 20, 150.	2.3	9
10	Relevance of the MHC region for breast cancer susceptibility in Asians. Breast Cancer, 2022, 29, 869-879.	1.3	1
11	Breast cancer risks associated with missense variants in breast cancer susceptibility genes. Genome Medicine, 2022, 14, 51.	3.6	19
12	Association between Breast Cancer Polygenic Risk Score and Chemotherapy-Induced Febrile Neutropenia: Null Results. Cancers, 2022, 14, 2714.	1.7	2
13	Epidemiological and ES cellâ€based functional evaluation of BRCA2 variants identified in families with breast cancer. Human Mutation, 2021, 42, 200-212.	1.1	4
14	Association between breast cancer risk and disease aggressiveness: Characterizing underlying gene expression patterns. International Journal of Cancer, 2021, 148, 884-894.	2.3	3
15	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
16	DNA methylation and breast cancer-associated variants. Breast Cancer Research and Treatment, 2021, 188, 713-727.	1.1	7
17	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
18	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

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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?. Cancers, 2021, 13, 2370.	1.7	4
20	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
21	Functionalized DMP-039 Hybrid Nanoparticle as a Novel mRNA Vector for Efficient Cancer Suicide Gene Therapy. International Journal of Nanomedicine, 2021, Volume 16, 5211-5232.	3.3	24
22	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
23	Genetic insights into biological mechanisms governing human ovarian ageing. Nature, 2021, 596, 393-397.	13.7	183
24	Single Micelle Vectors based on Lipid/Block Copolymer Compositions as mRNA Formulations for Efficient Cancer Immunogene Therapy. Molecular Pharmaceutics, 2021, 18, 4029-4045.	2.3	13
25	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
26	Breast cancer risk stratification for mammographic screening: A nationâ€wide screening cohort of 24,431 women in Singapore. Cancer Medicine, 2021, 10, 8182-8191.	1.3	6
27	Germline breast cancer susceptibility genes, tumor characteristics, and survival. Genome Medicine, 2021, 13, 185.	3.6	3
28	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. Nature Communications, 2020, 11, 3833.	5.8	88
29	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. American Journal of Human Genetics, 2020, 107, 837-848.	2.6	39
30	Treatment of Colon Cancer by Degradable rrPPC Nano-Conjugates Delivered STAT3 siRNA . International Journal of Nanomedicine, 2020, Volume 15, 9875-9890.	3.3	22
31	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. Scientific Reports, 2020, 10, 9688.	1.6	2
32	Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. Nature Communications, 2020, 11, 1217.	5.8	46
33	Impact of delayed treatment in women diagnosed with breast cancer: A populationâ€based study. Cancer Medicine, 2020, 9, 2435-2444.	1.3	46
34	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
35	Impact of deviation from guideline recommended treatment on breast cancer survival in Asia. Scientific Reports, 2020, 10, 1330.	1.6	14
36	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

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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. EBioMedicine, 2019, 48, 203-211.	2.7	14
38	Two truncating variants in FANCC and breast cancer risk. Scientific Reports, 2019, 9, 12524.	1.6	5
39	Joint association of mammographic density adjusted for age and body mass index and polygenic risk score with breast cancer risk. Breast Cancer Research, 2019, 21, 68.	2.2	31
40	The long-term prognostic and predictive capacity of cyclin D1 gene amplification in 2305 breast tumours. Breast Cancer Research, 2019, 21, 34.	2.2	48
41	Comparison of self-reported and register-based hospital medical data on comorbidities in women. Scientific Reports, 2019, 9, 3527.	1.6	13
42	Factors associated with false-positive mammography at first screen in an Asian population. PLoS ONE, 2019, 14, e0213615.	1.1	9
43	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
44	Prevalence of BRCA1 and BRCA2 pathogenic variants in a large, unselected breast cancer cohort. International Journal of Cancer, 2019, 144, 1195-1204.	2.3	31
45	A comprehensive tool for measuring mammographic density changes over time. Breast Cancer Research and Treatment, 2018, 169, 371-379.	1.1	45
46	Determinants of breast size in Asian women. Scientific Reports, 2018, 8, 1201.	1.6	19
47	Family History, Reproductive, and Lifestyle Risk Factors for Fibroadenoma and Breast Cancer. JNCI Cancer Spectrum, 2018, 2, pky051.	1.4	19
48	Differential Burden of Rare and Common Variants on Tumor Characteristics, Survival, and Mode of Detection in Breast Cancer. Cancer Research, 2018, 78, 6329-6338.	0.4	19
49	Inherited factors contribute to an inverse association between preeclampsia and breast cancer. Breast Cancer Research, 2018, 20, 6.	2.2	14
50	Common genetic variation and novel loci associated with volumetric mammographic density. Breast Cancer Research, 2018, 20, 30.	2.2	18
51	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. Nature Genetics, 2018, 50, 968-978.	9.4	184
52	BRCA2 Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. Cancer Research, 2017, 77, 2789-2799.	0.4	75
53	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. Nature Genetics, 2017, 49, 834-841.	9.4	426
54	Assessment of Breast Cancer Risk Factors Reveals Subtype Heterogeneity. Cancer Research, 2017, 77, 3708-3717.	0.4	87

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55	Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94.	13.7	1,099
56	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778.	9.4	289
57	Common shared genetic variation behind decreased risk of breast cancer in celiac disease. Scientific Reports, 2017, 7, 5942.	1.6	5
58	Longitudinal fluctuation in mammographic percent density differentiates between interval and screenâ€detected breast cancer. International Journal of Cancer, 2017, 140, 34-40.	2.3	6
59	Molecular Differences between Screen-Detected and Interval Breast Cancers Are Largely Explained by PAM50 Subtypes. Clinical Cancer Research, 2017, 23, 2584-2592.	3.2	15
60	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. Genetics in Medicine, 2017, 19, 599-603.	1.1	67
61	Associations between childhood body size and seventeen adverse outcomes: analysis of 65,057 European women. Scientific Reports, 2017, 7, 16917.	1.6	8
62	Body mass index and breast cancer survival: a Mendelian randomization analysis. International Journal of Epidemiology, 2017, 46, 1814-1822.	0.9	45
63	Risk and predictors of psoriasis in patients with breast cancer: a Swedish population-based cohort study. BMC Medicine, 2017, 15, 154.	2.3	13
64	Body size in early life and risk of breast cancer. Breast Cancer Research, 2017, 19, 84.	2.2	15
65	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. Breast Cancer Research, 2017, 19, 119.	2.2	43
66	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. PLoS Medicine, 2016, 13, e1002105.	3.9	118
67	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. PLoS ONE, 2016, 11, e0160316.	1.1	12
68	Fineâ€scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. International Journal of Cancer, 2016, 139, 1303-1317.	2.3	51
69	PALB2,CHEK2andATMrare variants and cancer risk: data from COGS. Journal of Medical Genetics, 2016, 53, 800-811.	1.5	174
70	Patient survival and tumor characteristics associated with CHEK2:p.I157T â€“ findings from the Breast Cancer Association Consortium. Breast Cancer Research, 2016, 18, 98.	2.2	39
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	Genetic predisposition to ductal carcinoma in situ of the breast. Breast Cancer Research, 2016, 18, 22.	2.2	43

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73	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
74	Five endometrial cancer risk loci identified through genome-wide association analysis. Nature Genetics, 2016, 48, 667-674.	9.4	77
75	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
76	Worse quality of life in young and recently diagnosed breast cancer survivors compared with female survivors of other cancers: A crossâ€sectional study. International Journal of Cancer, 2016, 139, 2415-2425.	2.3	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. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 1503-1510.	1.1	64
78	rs2735383, located at a microRNA binding site in the 3â€™UTR of NBS1, is not associated with breast cancer risk. Scientific Reports, 2016, 6, 36874.	1.6	2
79	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breastâ€“ovarian cancer susceptibility locus. Nature Communications, 2016, 7, 12675.	5.8	78
80	Common diseases as determinants of menopausal age. Human Reproduction, 2016, 31, 2856-2864.	0.4	42
81	Assessing within-woman changes in mammographic density: a comparison of fully versus semi-automated area-based approaches. Cancer Causes and Control, 2016, 27, 481-491.	0.8	15
82	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. Endocrine-Related Cancer, 2016, 23, 77-91.	1.6	62
83	Variants in 6q25.1 Are Associated with Mammographic Density in Malaysian Chinese Women. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 327-333.	1.1	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. Human Genetics, 2016, 135, 137-154.	1.8	8
85	Associations of Breast Cancer Risk Prediction Tools With Tumor Characteristics and Metastasis. Journal of Clinical Oncology, 2016, 34, 251-258.	0.8	45
86	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. Journal of the National Cancer Institute, 2016, 108, djv315.	3.0	77
87	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. Gynecologic Oncology, 2016, 141, 386-401.	0.6	18
88	RAD51B in Familial Breast Cancer. PLoS ONE, 2016, 11, e0153788.	1.1	26
89	A Combined Segmentation and Registration Framework for Bilateral and Temporal Mammogram Analysis. Journal of Medical Imaging and Health Informatics, 2016, 6, 380-386.	0.2	2
90	Investigation of geneâ€environment interactions between 47 newly identified breast cancer susceptibility loci and environmental risk factors. International Journal of Cancer, 2015, 136, E685-96.	2.3	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.	1.6	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.	1.1	6
93	Common germline polymorphisms associated with breast cancer-specific survival. Breast Cancer Research, 2015, 17, 58.	2.2	26
94	Identification of two novel mammographic density loci at 6Q25.1. Breast Cancer Research, 2015, 17, 75.	2.2	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.	2.2	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.	0.2	0
97	The Genetic Basis of Quality of Life in Healthy Swedish Women: A Candidate Gene Approach. PLoS ONE, 2015, 10, e0118292.	1.1	8
98	Mammographic Breast Density and Common Genetic Variants in Breast Cancer Risk Prediction. PLoS ONE, 2015, 10, e0136650.	1.1	20
99	SNP-SNP interaction analysis of NF-Î³B signaling pathway on breast cancer survival. Oncotarget, 2015, 6, 37979-37994.	0.8	20
100	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. Journal of the National Cancer Institute, 2015, 107, .	3.0	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.	4.1	59
102	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. Human Molecular Genetics, 2015, 24, 2966-2984.	1.4	40
103	Risk Factors and Tumor Characteristics of Interval Cancers by Mammographic Density. Journal of Clinical Oncology, 2015, 33, 1030-1037.	0.8	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.	2.6	76
105	Breast cancer genetic risk profile is differentially associated with interval and screen-detected breast cancers. Annals of Oncology, 2015, 26, 517-522.	0.6	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.	1.3	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.	9.4	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.	2.6	37

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109	Identification of Novel Genetic Markers of Breast Cancer Survival. Journal of the National Cancer Institute, 2015, 107, .	3.0	56
110	Candidate locus analysis of the TERTâ€“CLPTM1L cancer risk region on chromosome 5p15 identifies multiple independent variants associated with endometrial cancer risk. Human Genetics, 2015, 134, 231-245.	1.8	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.	2.2	20
112	Novel Associations between Common Breast Cancer Susceptibility Variants and Risk-Predicting Mammographic Density Measures. Cancer Research, 2015, 75, 2457-2467.	0.4	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.	9.4	357
114	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. Journal of the National Cancer Institute, 2015, 107, djv219.	3.0	99
115	Comprehensive genetic assessment of the ESR1 locus identifies a risk region for endometrial cancer. Endocrine-Related Cancer, 2015, 22, 851-861.	1.6	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.	1.1	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.	1.4	38
118	Fine-mapping of the HNF1B multicancer locus identifies candidate variants that mediate endometrial cancer risk. Human Molecular Genetics, 2015, 24, 1478-1492.	1.4	50
119	Ethnic Differences in Mammographic Densities: An Asian Cross-Sectional Study. PLoS ONE, 2015, 10, e0117568.	1.1	44
120	In Search for the Genetic Basis of Quality of Life in Healthy Swedish Womenâ€”A GWAS Study Using the iCOGS Custom Genotyping Array. PLoS ONE, 2015, 10, e0140563.	1.1	2
121	MicroRNA Related Polymorphisms and Breast Cancer Risk. PLoS ONE, 2014, 9, e109973.	1.1	49
122	Area and Volumetric Density Estimation in Processed Full-Field Digital Mammograms for Risk Assessment of Breast Cancer. PLoS ONE, 2014, 9, e110690.	1.1	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.	1.1	62
124	Digital mammographic density and breast cancer risk: a caseâ€“control study of six alternative density assessment methods. Breast Cancer Research, 2014, 16, 439.	2.2	165
125	Volumetric Mammographic Density: Heritability and Association With Breast Cancer Susceptibility Loci. Journal of the National Cancer Institute, 2014, 106, dju334-dju334.	3.0	21
126	Genetic Predisposition to In Situ and Invasive Lobular Carcinoma of the Breast. PLoS Genetics, 2014, 10, e1004285.	1.5	39

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127	Enhancement of Mammographic Density Measures in Breast Cancer Risk Prediction. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 1314-1323.	1.1	17
128	2q36.3 is associated with prognosis for oestrogen receptor-negative breast cancer patients treated with chemotherapy. Nature Communications, 2014, 5, 4051.	5.8	16
129	Polygenic scores associated with educational attainment in adults predict educational achievement and ADHD symptoms in children. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 510-520.	1.1	40
130	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
131	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
132	Automated breast tissue density assessment using high order regional texture descriptors in mammography. Proceedings of SPIE, 2014, , .	0.8	2
133	CYP2B6*6 is associated with increased breast cancer risk. International Journal of Cancer, 2014, 134, 426-430.	2.3	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. Human Molecular Genetics, 2014, 23, 1934-1946.	1.4	32
135	Identification of New Genetic Susceptibility Loci for Breast Cancer Through Consideration of Geneâ€Environment Interactions. Genetic Epidemiology, 2014, 38, 84-93.	0.6	28
136	Genome-wide association study identifies multiple loci associated with both mammographic density and breast cancer risk. Nature Communications, 2014, 5, 5303.	5.8	109
137	Mammographic Density Phenotypes and Risk of Breast Cancer: A Meta-analysis. Journal of the National Cancer Institute, 2014, 106, .	3.0	261
138	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
139	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
140	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. Nature Communications, 2014, 5, 4999.	5.8	105
141	Genetic variation in mitotic regulatory pathway genes is associated with breast tumor grade. Human Molecular Genetics, 2014, 23, 6034-6046.	1.4	12
142	The aetiology of convulsive status epilepticus: A study of 258 cases in Western China. Seizure: the Journal of the British Epilepsy Association, 2014, 23, 717-721.	0.9	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. American Journal of Human Genetics, 2013, 93, 1046-1060.	2.6	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. Nature Genetics, 2013, 45, 371-384.	9.4	493
146	Change of mammographic density predicts the risk of contralateral breast cancer - a case-control study. Breast Cancer Research, 2013, 15, R57.	2.2	44
147	Confirmation of the reduction of hormone replacement therapy-related breast cancer risk for carriers of the HSD17B1_937_G variant. Breast Cancer Research and Treatment, 2013, 138, 543-548.	1.1	10
148	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
149	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. Nature Genetics, 2013, 45, 353-361.	9.4	960
150	GWAS of 126,559 Individuals Identifies Genetic Variants Associated with Educational Attainment. Science, 2013, 340, 1467-1471.	6.0	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. Breast Cancer Research and Treatment, 2013, 138, 529-542.	1.1	18
152	Mammographic Density Reduction Is a Prognostic Marker of Response to Adjuvant Tamoxifen Therapy in Postmenopausal Patients With Breast Cancer. Journal of Clinical Oncology, 2013, 31, 2249-2256.	0.8	113
153	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
154	Calcium intake is not related to breast cancer risk among Singapore Chinese women. International Journal of Cancer, 2013, 133, 680-686.	2.3	15
155	Association of CYP2D6metabolizer status with mammographic density change in response to tamoxifen treatment. Breast Cancer Research, 2013, 15, R93.	2.2	10
156	The UGT1A6_19_GG genotype is a breast cancer risk factor. Frontiers in Genetics, 2013, 4, 104.	1.1	8
157	Common Breast Cancer Susceptibility Variants in LSP1 and RAD51L1 Are Associated with Mammographic Density Measures that Predict Breast Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 1156-1166.	1.1	101
158	Identification of a novel percent mammographic density locus at 12q24. Human Molecular Genetics, 2012, 21, 3299-3305.	1.4	31
159	Mammographic Breast Density and Breast Cancer: Evidence of a Shared Genetic Basis. Cancer Research, 2012, 72, 1478-1484.	0.4	54
160	CHEK2*1100delC Heterozygosity in Women With Breast Cancer Associated With Early Death, Breast Cancerâ€“Specific Death, and Increased Risk of a Second Breast Cancer. Journal of Clinical Oncology, 2012, 30, 4308-4316.	0.8	162
161	High-throughput mammographic-density measurement: a tool for risk prediction of breast cancer. Breast Cancer Research, 2012, 14, R114.	2.2	96
162	9q31.2-rs865686 as a Susceptibility Locus for Estrogen Receptor-Positive Breast Cancer: Evidence from the Breast Cancer Association Consortium. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 1783-1791.	1.1	17

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163	No evidence for association of inherited variation in genes involved in mitosis and percent mammographic density. Breast Cancer Research, 2012, 14, R7.	2.2	3
164	The postmenopausal hormone replacement therapy-related breast cancer risk is decreased in women carrying the CYP2C19*17 variant. Breast Cancer Research and Treatment, 2012, 131, 347-350.	1.1	6
165	Coffee consumption modifies risk of estrogen-receptor negative breast cancer. Breast Cancer Research, 2011, 13, R49.	2.2	46
166	Common variants in ZNF365 are associated with both mammographic density and breast cancer risk. Nature Genetics, 2011, 43, 185-187.	9.4	109
167	Genome-wide association study identifies a common variant associated with risk of endometrial cancer. Nature Genetics, 2011, 43, 451-454.	9.4	141
168	A combined analysis of genome-wide association studies in breast cancer. Breast Cancer Research and Treatment, 2011, 126, 717-727.	1.1	90
169	Blurring of High-Resolution Data Shows that the Effect of Intrinsic Nucleosome Occupancy on Transcription Factor Binding is Mostly Regional, Not Local. PLoS Computational Biology, 2010, 6, e1000649.	1.5	27
170	A genome-wide association scan on estrogen receptor-negative breast cancer. Breast Cancer Research, 2010, 12, R93.	2.2	35
171	Effects of childhood body size on breast cancer tumour characteristics. Breast Cancer Research, 2010, 12, R23.	2.2	13
172	Genetic variation in the estrogen metabolic pathway and mammographic density as an intermediate phenotype of breast cancer. Breast Cancer Research, 2010, 12, R19.	2.2	16