

Qin Wang

List of Publications by Year in descending order

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Version: 2024-02-01

172
papers

12,547
citations

44444

50
h-index

38517

99
g-index

180
all docs

180
docs citations

180
times ranked

18901
citing authors

#	ARTICLE	IF	CITATIONS
1	Rare germline copy number variants (CNVs) and breast cancer risk. <i>Communications Biology</i> , 2022, 5, 65.	2.0	6
2	Common variants in breast cancer risk loci predispose to distinct tumor subtypes. <i>Breast Cancer Research</i> , 2022, 24, 2.	2.2	15
3	Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes. <i>JAMA Oncology</i> , 2022, 8, e216744.	3.4	51
4	Characterising metabolomic signatures of lipid-modifying therapies through drug target mendelian randomisation. <i>PLoS Biology</i> , 2022, 20, e3001547.	2.6	69
5	Polygenic risk scores for prediction of breast cancer risk in Asian populations. <i>Genetics in Medicine</i> , 2022, 24, 586-600.	1.1	27
6	A Genome-Wide Gene-Based Gene-Environment Interaction Study of Breast Cancer in More than 90,000 Women. <i>Cancer Research Communications</i> , 2022, 2, 211-219.	0.7	6
7	Genome-wide and transcriptome-wide association studies of mammographic density phenotypes reveal novel loci. <i>Breast Cancer Research</i> , 2022, 24, 27.	2.2	15
8	A single-cell nanocoating of probiotics for enhanced amelioration of antibiotic-associated diarrhea. <i>Nature Communications</i> , 2022, 13, 2117.	5.8	74
9	Genome-wide interaction analysis of menopausal hormone therapy use and breast cancer risk among 62,370 women. <i>Scientific Reports</i> , 2022, 12, 6199.	1.6	2
10	Oral Administration of 5-Hydroxytryptophan Restores Gut Microbiota Dysbiosis in a Mouse Model of Depression. <i>Frontiers in Microbiology</i> , 2022, 13, 864571.	1.5	7
11	Polygenic risk scores for prediction of breast cancer risk in women of African ancestry: a cross-ancestry approach. <i>Human Molecular Genetics</i> , 2022, 31, 3133-3143.	1.4	11
12	Relevance of the MHC region for breast cancer susceptibility in Asians. <i>Breast Cancer</i> , 2022, 29, 869-879.	1.3	1
13	Breast cancer risks associated with missense variants in breast cancer susceptibility genes. <i>Genome Medicine</i> , 2022, 14, 51.	3.6	19
14	Distinct Reproductive Risk Profiles for Intrinsic-Like Breast Cancer Subtypes: Pooled Analysis of Population-Based Studies. <i>Journal of the National Cancer Institute</i> , 2022, 114, 1706-1719.	3.0	14
15	Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2021, 113, 329-337.	3.0	45
16	Metabolic profiling of angiotensin-like protein 3 and 4 inhibition: a drug-target Mendelian randomization analysis. <i>European Heart Journal</i> , 2021, 42, 1160-1169.	1.0	33
17	CYP3A7*1C allele: linking premenopausal oestrone and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , 2021, 124, 842-854.	2.9	5
18	A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. <i>Nature Communications</i> , 2021, 12, 1078.	5.8	19

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19	Breast Cancer Risk Genes " Association Analysis in More than 113,000 Women. <i>New England Journal of Medicine</i> , 2021, 384, 428-439.	13.9	532
20	Evaluating Polygenic Risk Scores for Breast Cancer in Women of African Ancestry. <i>Journal of the National Cancer Institute</i> , 2021, 113, 1168-1176.	3.0	41
21	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.	1.7	4
22	Cross-ancestry GWAS meta-analysis identifies six breast cancer loci in African and European ancestry women. <i>Nature Communications</i> , 2021, 12, 4198.	5.8	24
23	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.	2.6	6
24	Synthesis and Biological Evaluation of Falcarinol-Type Analogues as Potential Calcium Channel Blockers. <i>Journal of Natural Products</i> , 2021, 84, 2138-2148.	1.5	1
25	Association of germline genetic variants with breast cancer-specific survival in patient subgroups defined by clinic-pathological variables related to tumor biology and type of systemic treatment. <i>Breast Cancer Research</i> , 2021, 23, 86.	2.2	7
26	Mendelian randomisation study of smoking exposure in relation to breast cancer risk. <i>British Journal of Cancer</i> , 2021, 125, 1135-1145.	2.9	9
27	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	13.7	183
28	Interface-sensitized prodrug nanoaggregate as an effective in situ antitumor vaccine. <i>European Journal of Pharmaceutical Sciences</i> , 2021, 164, 105910.	1.9	4
29	Skin-inspired gelatin-based flexible bio-electronic hydrogel for wound healing promotion and motion sensing. <i>Biomaterials</i> , 2021, 276, 121026.	5.7	81
30	Synthesis, Anticancer Activity, Structure-Activity Relationship and Mechanistic Investigations of Falcarindiol Analogues. <i>ChemMedChem</i> , 2021, 16, 3569-3575.	1.6	2
31	Comparative Analyses of the Transcriptome and Proteome of <i>Escherichia coli</i> C321.Δ ^{3A} and Further Improving Its Noncanonical Amino Acids Containing Protein Expression Ability by Integration of T7 RNA Polymerase. <i>Frontiers in Microbiology</i> , 2021, 12, 744284.	1.5	5
32	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.	1.1	19
33	A crystalline covalent organic framework embedded with a crystalline supramolecular organic framework for efficient iodine capture. <i>Journal of Materials Chemistry A</i> , 2021, 9, 16961-16966.	5.2	43
34	Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. <i>Scientific Reports</i> , 2021, 11, 19787.	1.6	2
35	Enhancement of Activity and Thermostability of Keratinase From <i>Pseudomonas aeruginosa</i> CCTCC AB2013184 by Directed Evolution With Noncanonical Amino Acids. <i>Frontiers in Bioengineering and Biotechnology</i> , 2021, 9, 770907.	2.0	9
36	Effects of Nisin, Cecropin, and <i>Penthorum chinense</i> Pursh on the Intestinal Microbiome of Common Carp (<i>Cyprinus carpio</i>). <i>Frontiers in Nutrition</i> , 2021, 8, 729437.	1.6	17

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37	Dissection of the Functional Mechanism of Human Gut Bacterial Strain AD16 by Secondary Metabolitesâ€™ Identification, Network Pharmacology, and Experimental Validation. <i>Frontiers in Pharmacology</i> , 2021, 12, 706220.	1.6	0
38	Body mass index and type 2 diabetes and breast cancer survival: a Mendelian randomization study. <i>American Journal of Cancer Research</i> , 2021, 11, 3921-3934.	1.4	0
39	Iridoid, phenylethanoid and flavonoid glycosides from <i>Forsythia suspensa</i> . <i>Natural Product Research</i> , 2020, 34, 1320-1325.	1.0	12
40	Genetically Predicted Levels of DNA Methylation Biomarkers and Breast Cancer Risk: Data From 228â€™951 Women of European Descent. <i>Journal of the National Cancer Institute</i> , 2020, 112, 295-304.	3.0	35
41	Two new furofuran lignan glycoside derivatives from the fruit of <i>Forsythia suspensa</i> . <i>Journal of Asian Natural Products Research</i> , 2020, 22, 803-809.	0.7	2
42	Assessment of interactions between 205 breast cancer susceptibility loci and 13 established risk factors in relation to breast cancer risk, in the Breast Cancer Association Consortium. <i>International Journal of Epidemiology</i> , 2020, 49, 216-232.	0.9	21
43	Evaluation of associations between genetically predicted circulating protein biomarkers and breast cancer risk. <i>International Journal of Cancer</i> , 2020, 146, 2130-2138.	2.3	13
44	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
45	Middleâ€™and highâ€™molecular weight adiponectin levels in relation to nonalcoholic fatty liver disease. <i>Journal of Clinical Laboratory Analysis</i> , 2020, 34, e23148.	0.9	5
46	A Mendelian randomization analysis of circulating lipid traits and breast cancer risk. <i>International Journal of Epidemiology</i> , 2020, 49, 1117-1131.	0.9	41
47	Characterization of a thermostable phytase from <i>Bacillus licheniformis</i> WHU and further stabilization of the enzyme through disulfide bond engineering. <i>Enzyme and Microbial Technology</i> , 2020, 142, 109679.	1.6	15
48	Highly Diastereoselective Synthesis of 2,2-Disubstituted Cyclopentane-1,3-diols via Stepwise Ketone Reduction Enabling Concise Chirality Construction. <i>Journal of Organic Chemistry</i> , 2020, 85, 9599-9606.	1.7	4
49	Enhancement of keratin-degradation ability of the keratinase KerBL from <i>Bacillus licheniformis</i> WHU by proximity-triggered chemical crosslinking. <i>International Journal of Biological Macromolecules</i> , 2020, 163, 1458-1470.	3.6	12
50	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. <i>Nature Communications</i> , 2020, 11, 3833.	5.8	88
51	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. <i>Nature Genetics</i> , 2020, 52, 572-581.	9.4	265
52	Comparative analysis of the gut microbiota of grass carp fed with chicken faeces. <i>Environmental Science and Pollution Research</i> , 2020, 27, 32888-32898.	2.7	11
53	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , 2020, 10, 9688.	1.6	2
54	Insight Into the Effects of Nisin and Cecropin on the Oral Microbial Community of Rats by High-Throughput Sequencing. <i>Frontiers in Microbiology</i> , 2020, 11, 1082.	1.5	9

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55	Identification of novel breast cancer susceptibility loci in meta-analyses conducted among Asian and European descendants. <i>Nature Communications</i> , 2020, 11, 1217.	5.8	46
56	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	0.6	32
57	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312.	5.8	30
58	Prediction of contralateral breast cancer: external validation of risk calculators in 20 international cohorts. <i>Breast Cancer Research and Treatment</i> , 2020, 181, 423-434.	1.1	14
59	The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , 2019, 5, 38.	2.3	28
60	Excitation of One Fluorescent Probe at Two Different Wavelengths to Determine the Concentration and Enantiomeric Composition of Amino Acids. <i>Organic Letters</i> , 2019, 21, 9036-9039.	2.4	14
61	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.	2.7	14
62	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , 2019, 9, 12524.	1.6	5
63	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88
64	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.	2.2	31
65	A combination of the immunohistochemical markers CK7 and SATB2 is highly sensitive and specific for distinguishing primary ovarian mucinous tumors from colorectal and appendiceal metastases. <i>Modern Pathology</i> , 2019, 32, 1834-1846.	2.9	54
66	Authors' response: Associations of obesity and circulating insulin and glucose with breast cancer risk. <i>International Journal of Epidemiology</i> , 2019, 48, 1016-1017.	0.9	1
67	The functional ALDH2 polymorphism is associated with breast cancer risk: A pooled analysis from the Breast Cancer Association Consortium. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e707.	0.6	9
68	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	5.8	90
69	PLCE1 Polymorphisms and Risk of Esophageal and Gastric Cancer in a Northwestern Chinese Population. <i>BioMed Research International</i> , 2019, 2019, 1-10.	0.9	7
70	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019, 120, 647-657.	2.9	52
71	Prediction and clinical utility of a contralateral breast cancer risk model. <i>Breast Cancer Research</i> , 2019, 21, 144.	2.2	24
72	NLRC5 mediates cell proliferation, migration, and invasion by regulating the Wnt/ β -catenin signalling pathway in clear cell renal cell carcinoma. <i>Cancer Letters</i> , 2019, 444, 9-19.	3.2	34

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73	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. American Journal of Human Genetics, 2019, 104, 21-34.	2.6	711
74	Blockade of YAP alleviates hepatic fibrosis through accelerating apoptosis and reversion of activated hepatic stellate cells. Molecular Immunology, 2019, 107, 29-40.	1.0	63
75	Targeted Resequencing of the Coding Sequence of 38 Genes Near Breast Cancer GWAS Loci in a Large Case-Control Study. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 822-825.	1.1	7
76	Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis. International Journal of Epidemiology, 2019, 48, 795-806.	0.9	81
77	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
78	Emerging roles of hsa_circ_0005075 targeting miR-431 in the progress of HCC. Biomedicine and Pharmacotherapy, 2018, 99, 848-858.	2.5	43
79	Etiology of hormone receptor positive breast cancer differs by levels of histologic grade and proliferation. International Journal of Cancer, 2018, 143, 746-757.	2.3	19
80	Genetic overlap between endometriosis and endometrial cancer: evidence from cross-disease genetic correlation and GWAS meta-analyses. Cancer Medicine, 2018, 7, 1978-1987.	1.3	62
81	Joint associations of a polygenic risk score and environmental risk factors for breast cancer in the Breast Cancer Association Consortium. International Journal of Epidemiology, 2018, 47, 526-536.	0.9	88
82	SENP2 alleviates CCl4-induced liver fibrosis by promoting activated hepatic stellate cell apoptosis and reversion. Toxicology Letters, 2018, 289, 86-98.	0.4	22
83	Enhanced Turnover for the P450 119 Peroxygenase-Catalyzed Asymmetric Epoxidation of Styrenes by Random Mutagenesis. Chemistry - A European Journal, 2018, 24, 2741-2749.	1.7	16
84	Alkylation of 3-FormylBINOL and Its Strong Effect on the Fluorescence Recognition of 1,3-Diaminopropane. European Journal of Organic Chemistry, 2018, 2018, 4972-4977.	1.2	9
85	Association of p16 expression with prognosis varies across ovarian carcinoma histotypes: an Ovarian Tumor Tissue Analysis consortium study. Journal of Pathology: Clinical Research, 2018, 4, 250-261.	1.3	70
86	The Long Non-coding RNA MEG3/miR-let-7c-5p Axis Regulates Ethanol-Induced Hepatic Steatosis and Apoptosis by Targeting NLRC5. Frontiers in Pharmacology, 2018, 9, 302.	1.6	30
87	DNA Methylation of PTCIS Enhances Hepatic Stellate Cells Activation and Liver Fibrogenesis. Frontiers in Pharmacology, 2018, 9, 553.	1.6	23
88	Highly Enantioselective Synthesis and Anticancer Activities of Chiral Conjugated Diynols. ChemBioChem, 2018, 19, 2293-2299.	1.3	6
89	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
90	Anti-inflammatory and neuroprotective effects of sanguinarine following cerebral ischemia in rats. Experimental and Therapeutic Medicine, 2017, 13, 263-268.	0.8	38

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91	<i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017, 77, 2789-2799.	0.4	75
92	Insights into the molecular mechanisms of <i>Polygonum multiflorum</i> Thunb-induced liver injury: a computational systems toxicology approach. <i>Acta Pharmacologica Sinica</i> , 2017, 38, 719-732.	2.8	30
93	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.	9.4	426
94	Evaluating genetic variants associated with breast cancer risk in high and moderate-penetrance genes in Asians. <i>Carcinogenesis</i> , 2017, 38, 511-518.	1.3	38
95	In silico prediction of serious eye irritation or corrosion potential of chemicals. <i>RSC Advances</i> , 2017, 7, 6697-6703.	1.7	37
96	In silico prediction of chemical subcellular localization via multi-classification methods. <i>MedChemComm</i> , 2017, 8, 1225-1234.	3.5	15
97	Access to Fluorescent Azines from Heterocyclic Carbene Precursors and <i>N</i> -Tosylhydrazones. <i>Advanced Synthesis and Catalysis</i> , 2017, 359, 1825-1830.	2.1	5
98	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099
99	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
100	Gene-environment interactions involving functional variants: Results from the Breast Cancer Association Consortium. <i>International Journal of Cancer</i> , 2017, 141, 1830-1840.	2.3	20
101	Rare, protein-truncating variants in <i>ATM</i> , <i>CHEK2</i> and <i>PALB2</i> , but not <i>XRCC2</i> , are associated with increased breast cancer risks. <i>Journal of Medical Genetics</i> , 2017, 54, 732-741.	1.5	68
102	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017, 19, 599-603.	1.1	67
103	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017, 46, 1814-1822.	0.9	45
104	TP53-based interaction analysis identifies cis-eQTL variants for TP53BP2, FBXO28, and FAM53A that associate with survival and treatment outcome in breast cancer. <i>Oncotarget</i> , 2017, 8, 18381-18398.	0.8	14
105	<i>PHIP</i> - a novel candidate breast cancer susceptibility locus on 6q14.1. <i>Oncotarget</i> , 2017, 8, 102769-102782.	0.8	9
106	Unraveling Molecular Differences of Gastric Cancer by Label-Free Quantitative Proteomics Analysis. <i>International Journal of Molecular Sciences</i> , 2016, 17, 69.	1.8	31
107	Association of breast cancer risk with genetic variants showing differential allelic expression: Identification of a novel breast cancer susceptibility locus at 4q21. <i>Oncotarget</i> , 2016, 7, 80140-80163.	0.8	31
108	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.	3.9	118

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109	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. PLoS ONE, 2016, 11, e0160316.	1.1	12
110	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
111	Combined administration of propranolol and AG490 offers better effects on portal hypertensive rats with cirrhosis. Journal of Gastroenterology and Hepatology (Australia), 2016, 31, 1037-1044.	1.4	15
112	Activation of innate antiviral immune response via double-stranded RNA-dependent RLR receptor-mediated necroptosis. Scientific Reports, 2016, 6, 22550.	1.6	12
113	Patient survival and tumor characteristics associated with CHEK2:p.1157T findings from the Breast Cancer Association Consortium. Breast Cancer Research, 2016, 18, 98.	2.2	39
114	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
115	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
116	Genetic predisposition to ductal carcinoma in situ of the breast. Breast Cancer Research, 2016, 18, 22.	2.2	43
117	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
118	Five endometrial cancer risk loci identified through genome-wide association analysis. Nature Genetics, 2016, 48, 667-674.	9.4	77
119	Effects of hormonal contraception on systemic metabolism: cross-sectional and longitudinal evidence. International Journal of Epidemiology, 2016, 45, 1445-1457.	0.9	62
120	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
121	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
122	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
123	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
124	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. Cancer Discovery, 2016, 6, 1052-1067.	7.7	157
125	Genome-wide study for circulating metabolites identifies 62 loci and reveals novel systemic effects of LPA. Nature Communications, 2016, 7, 11122.	5.8	576
126	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. Nature Communications, 2016, 7, 12675.	5.8	78

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127	Fine scale mapping of the 17q22 breast cancer locus using dense SNPs, genotyped within the Collaborative Oncological Gene-Environment Study (COGs). <i>Scientific Reports</i> , 2016, 6, 32512.	1.6	19
128	Metabolic profiling of alcohol consumption in 9778 young adults. <i>International Journal of Epidemiology</i> , 2016, 45, 1493-1506.	0.9	90
129	Anticancer effects of morin-7-sulphate sodium, a flavonoid derivative, in mouse melanoma cells. <i>Biomedicine and Pharmacotherapy</i> , 2016, 84, 909-916.	2.5	27
130	Prognostic value of automated Ki67 scoring in breast cancer: a centralised evaluation of 8088 patients from 10 study groups. <i>Breast Cancer Research</i> , 2016, 18, 104.	2.2	56
131	Metabolic profiling of pregnancy: cross-sectional and longitudinal evidence. <i>BMC Medicine</i> , 2016, 14, 205.	2.3	150
132	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2016, 23, 77-91.	1.6	62
133	High-throughput automated scoring of Ki67 in breast cancer tissue microarrays from the Breast Cancer Association Consortium. <i>Journal of Pathology: Clinical Research</i> , 2016, 2, 138-153.	1.3	19
134	Combined genetic and splicing analysis of BRCA1 c.[594-2A>C; 641A>G] highlights the relevance of naturally occurring in-frame transcripts for developing disease gene variant classification algorithms. <i>Human Molecular Genetics</i> , 2016, 25, 2256-2268.	1.4	106
135	Alu RNA accumulation in hyperglycemia augments oxidative stress and impairs eNOS and SOD2 expression in endothelial cells. <i>Molecular and Cellular Endocrinology</i> , 2016, 426, 91-100.	1.6	27
136	No evidence that protein truncating variants in <i>BRIP1</i> are associated with breast cancer risk: implications for gene panel testing. <i>Journal of Medical Genetics</i> , 2016, 53, 298-309.	1.5	94
137	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016, 48, 374-386.	9.4	125
138	Chemical profiling of Qixue Shuangbu Tincture by ultra-performance liquid chromatography with electrospray ionization quadrupole-time-of-flight high-definition mass spectrometry (UPLC-QTOF/MS). <i>Chinese Journal of Natural Medicines</i> , 2016, 14, 141-146.	0.7	15
139	Metabolomic Profiling of Statin Use and Genetic Inhibition of HMG-CoA Reductase. <i>Journal of the American College of Cardiology</i> , 2016, 67, 1200-1210.	1.2	173
140	Heterogeneity of luminal breast cancer characterised by immunohistochemical expression of basal markers. <i>British Journal of Cancer</i> , 2016, 114, 298-304.	2.9	7
141	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.	1.8	8
142	RAD51B in Familial Breast Cancer. <i>PLoS ONE</i> , 2016, 11, e0153788.	1.1	26
143	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. <i>Scientific Reports</i> , 2015, 5, 17369.	1.6	35
144	A polymorphism in the base excision repair gene PARP2 is associated with differential prognosis by chemotherapy among postmenopausal breast cancer patients. <i>BMC Cancer</i> , 2015, 15, 978.	1.1	6

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145	Common germline polymorphisms associated with breast cancer-specific survival. <i>Breast Cancer Research</i> , 2015, 17, 58.	2.2	26
146	A comprehensive evaluation of interaction between genetic variants and use of menopausal hormone therapy on mammographic density. <i>Breast Cancer Research</i> , 2015, 17, 110.	2.2	19
147	Development of a HPLC Method for the Quantitative Determination of Capsaicin in Collagen Sponge. <i>International Journal of Analytical Chemistry</i> , 2015, 2015, 1-6.	0.4	19
148	SNP-SNP interaction analysis of NF- κ B signaling pathway on breast cancer survival. <i>Oncotarget</i> , 2015, 6, 37979-37994.	0.8	20
149	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	3.0	428
150	Sex hormone-binding globulin associations with circulating lipids and metabolites and the risk for type 2 diabetes: observational and causal effect estimates. <i>International Journal of Epidemiology</i> , 2015, 44, 623-637.	0.9	83
151	Crowdsourcing the General Public for Large Scale Molecular Pathology Studies in Cancer. <i>EBioMedicine</i> , 2015, 2, 681-689.	2.7	56
152	Fine-Scale Mapping of the 5q11.2 Breast Cancer Locus Reveals at Least Three Independent Risk Variants Regulating MAP3K1. <i>American Journal of Human Genetics</i> , 2015, 96, 5-20.	2.6	76
153	Metabolite Profiling and Cardiovascular Event Risk. <i>Circulation</i> , 2015, 131, 774-785.	1.6	547
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