

Anthony J Swerdlow

List of Publications by Year in Descending Order

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

193
papers

11,724
citations

54
h-index

104
g-index

206
ext. papers

14,946
ext. citations

11.1
avg, IF

5.07
L-index

#	Paper	IF	Citations
193	Rare germline copy number variants (CNVs) and breast cancer risk.. <i>Communications Biology</i> , 2022 , 5, 65	6.7	0
192	Common variants in breast cancer risk loci predispose to distinct tumor subtypes.. <i>Breast Cancer Research</i> , 2022 , 24, 2	8.3	3
191	Risk of breast cancer in men in relation to weight change: a national case-control study in England and Wales.. <i>International Journal of Cancer</i> , 2022 ,	7.5	1
190	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
189	Genome-wide and transcriptome-wide association studies of mammographic density phenotypes reveal novel loci.. <i>Breast Cancer Research</i> , 2022 , 24, 27	8.3	1
188	Genome-wide interaction analysis of menopausal hormone therapy use and breast cancer risk among 62,370 women.. <i>Scientific Reports</i> , 2022 , 12, 6199	4.9	
187	Infertility and risk of breast cancer in men: a national case-control study in England and Wales.. <i>Breast Cancer Research</i> , 2022 , 24, 29	8.3	
186	Cross-Cancer Genome-Wide Association Study of Endometrial Cancer and Epithelial Ovarian Cancer Identifies Genetic Risk Regions Associated with Risk of Both Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021 , 30, 217-228	4	7
185	Validated biomarker assays confirm ARID1A loss is confounded with MMR deficiency, CD8 TIL infiltration, and provides no independent prognostic value in endometriosis-associated ovarian carcinomas.. <i>Journal of Pathology</i> , 2021 ,	9.4	3
184	Risk of thyroid disorders in adult and childhood Hodgkin lymphoma survivors 40 years after treatment. <i>Leukemia and Lymphoma</i> , 2021 , 1-11	1.9	
183	MCM3 is a novel proliferation marker associated with longer survival for patients with tubo-ovarian high-grade serous carcinoma. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2021 ,	5.1	2
182	Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. <i>Scientific Reports</i> , 2021 , 11, 19787	4.9	0
181	Functional antibody and T cell immunity following SARS-CoV-2 infection, including by variants of concern, in patients with cancer: the CAPTURE study.. <i>Nature Cancer</i> , 2021 , 2, 1321-1337	15.4	17
180	Prospective evaluation of a breast-cancer risk model integrating classical risk factors and polygenic risk in 15 cohorts from six countries. <i>International Journal of Epidemiology</i> , 2021 ,	7.8	6
179	Breast Cancer Risk Factors and Circulating Anti-Müllerian Hormone Concentration in Healthy Premenopausal Women. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021 , 106, e4542-e4553	5.6	
178	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	11	1
177	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	9.7	14

176	Common Susceptibility Loci for Male Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2021 , 113, 453-461	9.7	4
175	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	8.7	2
174	Comparative validation of the BOADICEA and Tyrer-Cuzick breast cancer risk models incorporating classical risk factors and polygenic risk in a population-based prospective cohort of women of European ancestry. <i>Breast Cancer Research</i> , 2021 , 23, 22	8.3	12
173	Obesity and Breast Cancer Risk in Men: A National Case-Control Study in England and Wales. <i>JNCI Cancer Spectrum</i> , 2021 , 5, pkab078	4.6	2
172	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	8.3	1
171	Mendelian randomisation study of smoking exposure in relation to breast cancer risk. <i>British Journal of Cancer</i> , 2021 , 125, 1135-1145	8.7	0
170	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021 , 596, 393-397	10.4	28
169	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	36.3	76
168	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , 2020 , 10, 9688	4.9	2
167	The Risk of Ovarian Cancer Increases with an Increase in the Lifetime Number of Ovulatory Cycles: An Analysis from the Ovarian Cancer Cohort Consortium (OC3). <i>Cancer Research</i> , 2020 , 80, 1210-1218	10.1	17
166	Transcriptome-wide association study of breast cancer risk by estrogen-receptor status. <i>Genetic Epidemiology</i> , 2020 , 44, 442-468	2.6	9
165	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020 , 11, 312	17.4	20
164	Adult weight change and premenopausal breast cancer risk: A prospective pooled analysis of data from 628,463 women. <i>International Journal of Cancer</i> , 2020 , 147, 1306-1314	7.5	8
163	Genome-wide association meta-analyses combining multiple risk phenotypes provide insights into the genetic architecture of cutaneous melanoma susceptibility. <i>Nature Genetics</i> , 2020 , 52, 494-504	36.3	39
162	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020 , 52, 56-73	36.3	56
161	Cancer, COVID-19, and Antiviral Immunity: The CAPTURE Study. <i>Cell</i> , 2020 , 183, 4-10	56.2	23
160	Long-term mortality after childhood growth hormone treatment: the SAGhE cohort study. <i>Lancet Diabetes and Endocrinology</i> , 2020 , 8, 683-692	18.1	22
159	Comparative Validation of Breast Cancer Risk Prediction Models and Projections for Future Risk Stratification. <i>Journal of the National Cancer Institute</i> , 2020 , 112, 278-285	9.7	36

158	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , 2019 , 9, 12524	4.9	2
157	Reply to Comment on: "Night shift work and risk of breast cancer in women: the Generations Study cohort". <i>British Journal of Cancer</i> , 2019 , 121, 723-724	8.7	
156	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019 , 10, 431	17.4	45
155	Risk of Meningioma in European Patients Treated With Growth Hormone in Childhood: Results From the SAGhE Cohort. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2019 , 104, 658-664	5.6	17
154	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	9.8	21
153	Night shift work and risk of breast cancer in women: the Generations Study cohort. <i>British Journal of Cancer</i> , 2019 , 121, 172-179	8.7	26
152	Association analyses identify 31 new risk loci for colorectal cancer susceptibility. <i>Nature Communications</i> , 2019 , 10, 2154	17.4	81
151	Breast cancer risk prediction in women aged 35-50 years: impact of including sex hormone concentrations in the Gail model. <i>Breast Cancer Research</i> , 2019 , 21, 42	8.3	16
150	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019 , 10, 1741	17.4	47
149	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019 , 120, 647-657	8.7	28
148	Analgesic Use and Ovarian Cancer Risk: An Analysis in the Ovarian Cancer Cohort Consortium. <i>Journal of the National Cancer Institute</i> , 2019 , 111, 137-145	9.7	28
147	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	7.8	16
146	The :p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , 2019 , 5, 38	7.8	12
145	Data must be shared-also with researchers outside of Europe. <i>Lancet, The</i> , 2019 , 394, 1902-1903	40	3
144	Transcriptome-Wide Association Study Identifies New Candidate Susceptibility Genes for Glioma. <i>Cancer Research</i> , 2019 , 79, 2065-2071	10.1	16
143	Mortality and cancer incidence in carriers of constitutional t(11;22)(q23;q11) translocations: A prospective study. <i>International Journal of Cancer</i> , 2019 , 145, 1493-1498	7.5	2
142	Sharing data safely while preserving privacy. <i>Lancet, The</i> , 2019 , 394, 1902	40	3
141	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019 , 104, 21-34	11	363

140	Genetic susceptibility to radiation-induced breast cancer after Hodgkin lymphoma. <i>Blood</i> , 2019 , 133, 1130-1139	2.2	17
139	Genome-wide association study of anti-Müllerian hormone levels in pre-menopausal women of late reproductive age and relationship with genetic determinants of reproductive lifespan. <i>Human Molecular Genetics</i> , 2019 , 28, 1392-1401	5.6	9
138	Response to "Comment on "Domestic light at night and breast cancer risk: a prospective analysis of 105 000 UK women in the Generations Study". <i>British Journal of Cancer</i> , 2019 , 120, 278	8.7	
137	Mortality and Cancer Incidence in Carriers of Balanced Robertsonian Translocations: A National Cohort Study. <i>American Journal of Epidemiology</i> , 2019 , 188, 500-508	3.8	4
136	Genetic Data from Nearly 63,000 Women of European Descent Predicts DNA Methylation Biomarkers and Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , 2019 , 79, 505-517	10.1	28
135	The BRCA2 c.68-7T>A variant is not pathogenic: A model for clinical calibration of spliceogenicity. <i>Human Mutation</i> , 2018 , 39, 729-741	4.7	16
134	Breast cancer risk in relation to history of preeclampsia and hyperemesis gravidarum: Prospective analysis in the Generations Study. <i>International Journal of Cancer</i> , 2018 , 143, 782-792	7.5	8
133	Etiology of hormone receptor positive breast cancer differs by levels of histologic grade and proliferation. <i>International Journal of Cancer</i> , 2018 , 143, 746-757	7.5	9
132	Genetic overlap between endometriosis and endometrial cancer: evidence from cross-disease genetic correlation and GWAS meta-analyses. <i>Cancer Medicine</i> , 2018 , 7, 1978-1987	4.8	40
131	Genome-wide association study identifies susceptibility loci for B-cell childhood acute lymphoblastic leukemia. <i>Nature Communications</i> , 2018 , 9, 1340	17.4	39
130	Domestic light at night and breast cancer risk: a prospective analysis of 105 000 UK women in the Generations Study. <i>British Journal of Cancer</i> , 2018 , 118, 600-606	8.7	30
129	Joint associations of a polygenic risk score and environmental risk factors for breast cancer in the Breast Cancer Association Consortium. <i>International Journal of Epidemiology</i> , 2018 , 47, 526-536	7.8	53
128	Circulating anti-Müllerian hormone and breast cancer risk: A study in ten prospective cohorts. <i>International Journal of Cancer</i> , 2018 , 142, 2215-2226	7.5	20
127	E-cadherin breast tumor expression, risk factors and survival: Pooled analysis of 5,933 cases from 12 studies in the Breast Cancer Association Consortium. <i>Scientific Reports</i> , 2018 , 8, 6574	4.9	19
126	Influence of obesity-related risk factors in the aetiology of glioma. <i>British Journal of Cancer</i> , 2018 , 118, 1020-1027	8.7	22
125	Effect of population breast screening on breast cancer mortality to 2005 in England and Wales: A nested case-control study within a cohort of one million women. <i>Journal of Medical Screening</i> , 2018 , 25, 76-81	1.4	4
124	Association of Body Mass Index and Age With Subsequent Breast Cancer Risk in Premenopausal Women. <i>JAMA Oncology</i> , 2018 , 4, e181771	13.4	129
123	Association of p16 expression with prognosis varies across ovarian carcinoma histotypes: an Ovarian Tumor Tissue Analysis consortium study. <i>Journal of Pathology: Clinical Research</i> , 2018 , 4, 250-261	5.3	38

122	A Transcriptome-Wide Association Study Among 97,898 Women to Identify Candidate Susceptibility Genes for Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , 2018 , 78, 5419-5430	10.1	32
121	The National Cancer Institute Cohort Consortium: An International Pooling Collaboration of 58 Cohorts from 20 Countries. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018 , 27, 1307-1319	4	11
120	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018 , 9, 3166	17.4	70
119	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	36.3	101
118	Maternal breast cancer risk in relation to birthweight and gestation of her offspring. <i>Breast Cancer Research</i> , 2018 , 20, 110	8.3	2
117	Response to Comment on "Domestic light at night and breast cancer risk: a prospective analysis of 105 000 UK women in the Generations Study". <i>British Journal of Cancer</i> , 2018 , 119, 1170	8.7	
116	Identification of multiple risk loci and regulatory mechanisms influencing susceptibility to multiple myeloma. <i>Nature Communications</i> , 2018 , 9, 3707	17.4	57
115	Genome-wide association study implicates immune dysfunction in the development of Hodgkin lymphoma. <i>Blood</i> , 2018 , 132, 2040-2052	2.2	10
114	Reply to Comment on "Domestic light at night and breast cancer risk: a prospective analysis of 105 000 UK women in the Generations Study". <i>British Journal of Cancer</i> , 2018 , 118, 1537	8.7	
113	Genome-wide association analysis identifies a meningioma risk locus at 11p15.5. <i>Neuro-Oncology</i> , 2018 , 20, 1485-1493	1	13
112	Genome-wide association analysis of chronic lymphocytic leukaemia, Hodgkin lymphoma and multiple myeloma identifies pleiotropic risk loci. <i>Scientific Reports</i> , 2017 , 7, 41071	4.9	27
111	Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017 , 77, 2789-2799	10.1	49
110	Cancer Risks in Patients Treated With Growth Hormone in Childhood: The SAGhE European Cohort Study. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2017 , 102, 1661-1672	5.6	79
109	The Premenopausal Breast Cancer Collaboration: A Pooling Project of Studies Participating in the National Cancer Institute Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017 , 26, 1360-1369	4	16
108	Genome-wide association study of glioma subtypes identifies specific differences in genetic susceptibility to glioblastoma and non-glioblastoma tumors. <i>Nature Genetics</i> , 2017 , 49, 789-794	36.3	163
107	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017 , 49, 680-691	36.3	190
106	Effect of population breast screening on breast cancer mortality up to 2005 in England and Wales: an individual-level cohort study. <i>British Journal of Cancer</i> , 2017 , 116, 246-252	8.7	30
105	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017 , 551, 92-94	50.4	643

104	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017 , 49, 1767-1778	36.3	186
103	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. <i>Breast Cancer Research</i> , 2017 , 19, 119	8.3	26
102	Smoking and risk of breast cancer in the Generations Study cohort. <i>Breast Cancer Research</i> , 2017 , 19, 118	8.3	57
101	Gene-environment interactions involving functional variants: Results from the Breast Cancer Association Consortium. <i>International Journal of Cancer</i> , 2017 , 141, 1830-1840	7.5	13
100	Genome-wide association study of classical Hodgkin lymphoma identifies key regulators of disease susceptibility. <i>Nature Communications</i> , 2017 , 8, 1892	17.4	24
99	Retrospective methods to estimate radiation dose at the site of breast cancer development after Hodgkin lymphoma radiotherapy. <i>Clinical and Translational Radiation Oncology</i> , 2017 , 7, 20-27	4.6	4
98	Family history and risk of breast cancer: an analysis accounting for family structure. <i>Breast Cancer Research and Treatment</i> , 2017 , 165, 193-200	4.4	86
97	Childhood body size and pubertal timing in relation to adult mammographic density phenotype. <i>Breast Cancer Research</i> , 2017 , 19, 13	8.3	14
96	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017 , 19, 599-603	6.3	51
95	- a novel candidate breast cancer susceptibility locus on 6q14.1. <i>Oncotarget</i> , 2017 , 8, 102769-102782	3.3	3
94	An intergenic risk locus containing an enhancer deletion in 2q35 modulates breast cancer risk by deregulating IGFBP5 expression. <i>Human Molecular Genetics</i> , 2016 , 25, 3863-3876	5.6	24
93	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	4	42
92	Psychological stress, adverse life events and breast cancer incidence: a cohort investigation in 106,000 women in the United Kingdom. <i>Breast Cancer Research</i> , 2016 , 18, 72	8.3	31
91	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. <i>Cancer Discovery</i> , 2016 , 6, 1052-674.4	24.4	104
90	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016 , 7, 11375	17.4	64
89	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016 , 7, 12675	17.4	53
88	The Intracranial Distribution of Gliomas in Relation to Exposure From Mobile Phones: Analyses From the INTERPHONE Study. <i>American Journal of Epidemiology</i> , 2016 , 184, 818-828	3.8	19
87	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	8.3	44

86	Age- and Tumor Subtype-Specific Breast Cancer Risk Estimates for CHEK2*1100delC Carriers. <i>Journal of Clinical Oncology</i> , 2016 , 34, 2750-60	2.2	107
85	Ovarian Cancer Risk Factors by Histologic Subtype: An Analysis From the Ovarian Cancer Cohort Consortium. <i>Journal of Clinical Oncology</i> , 2016 , 34, 2888-98	2.2	236
84	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2016 , 23, 77-91	5.7	41
83	No evidence that protein truncating variants in BRIP1 are associated with breast cancer risk: implications for gene panel testing. <i>Journal of Medical Genetics</i> , 2016 , 53, 298-309	5.8	83
82	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016 , 48, 374-86	36.3	93
81	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, 137-54	6.3	6
80	RAD51B in Familial Breast Cancer. <i>PLoS ONE</i> , 2016 , 11, e0153788	3.7	18
79	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	3.3	21
78	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	11.6	80
77	Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. <i>International Journal of Cancer</i> , 2016 , 139, 1303-1317	7.5	26
76	PALB2, CHEK2 and ATM rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016 , 53, 800-811	5.8	121
75	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	8.3	25
74	Genetic predisposition to ductal carcinoma in situ of the breast. <i>Breast Cancer Research</i> , 2016 , 18, 22	8.3	31
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, 679-93	2.8	15
72	Five endometrial cancer risk loci identified through genome-wide association analysis. <i>Nature Genetics</i> , 2016 , 48, 667-674	36.3	56
71	Length of FMR1 repeat alleles within the normal range does not substantially affect the risk of early menopause. <i>Human Reproduction</i> , 2016 , 31, 2396-403	5.7	7
70	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	11	43
69	Menopausal hormone therapy and breast cancer: what is the true size of the increased risk?. <i>British Journal of Cancer</i> , 2016 , 115, 607-15	8.7	47

68	Inherited variants in the inner centromere protein (INCENP) gene of the chromosomal passenger complex contribute to the susceptibility of ER-negative breast cancer. <i>Carcinogenesis</i> , 2015 , 36, 256-71	4.6	12
67	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015 , 47, 373-80	36.3	406
66	Description of the SAGHe Cohort: A Large European Study of Mortality and Cancer Incidence Risks after Childhood Treatment with Recombinant Growth Hormone. <i>Hormone Research in Paediatrics</i> , 2015 , 84, 172-83	3.3	32
65	Genome-wide meta-analysis identifies five new susceptibility loci for cutaneous malignant melanoma. <i>Nature Genetics</i> , 2015 , 47, 987-995	36.3	162
64	Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression. <i>American Journal of Human Genetics</i> , 2015 , 97, 22-34	11	26
63	Candidate locus analysis of the TERT-CLPTM1L cancer risk region on chromosome 5p15 identifies multiple independent variants associated with endometrial cancer risk. <i>Human Genetics</i> , 2015 , 134, 231-45	6.3	30
62	Genome-wide association study identifies multiple susceptibility loci for glioma. <i>Nature Communications</i> , 2015 , 6, 8559	17.4	81
61	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. <i>Nature Genetics</i> , 2015 , 47, 1294-1303	36.3	226
60	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. <i>Journal of the National Cancer Institute</i> , 2015 , 107,	9.7	74
59	Comprehensive genetic assessment of the ESR1 locus identifies a risk region for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2015 , 22, 851-61	5.7	19
58	Fine-scale mapping of the 4q24 locus identifies two independent loci associated with breast cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 1680-91	4	17
57	Tobacco and alcohol in relation to male breast cancer: an analysis of the male breast cancer pooling project consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 520-31	4	15
56	Fine-mapping of the HNF1B multicancer locus identifies candidate variants that mediate endometrial cancer risk. <i>Human Molecular Genetics</i> , 2015 , 24, 1478-92	5.6	46
55	Temporal stability and determinants of white blood cell DNA methylation in the breakthrough generations study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 221-9	4	43
54	Genome-wide homozygosity signature and risk of Hodgkin lymphoma. <i>Scientific Reports</i> , 2015 , 5, 14315	4.9	12
53	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	4.9	27
52	Prediction of breast cancer risk based on profiling with common genetic variants. <i>Journal of the National Cancer Institute</i> , 2015 , 107,	9.7	324
51	Mitochondrial DNA Copy Number in Peripheral Blood Cells and Risk of Developing Breast Cancer. <i>Cancer Research</i> , 2015 , 75, 2844-50	10.1	32

50	Development and validation of a melanoma risk score based on pooled data from 16 case-control studies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 817-24	4	18
49	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , 2015 , 24, 2966-84	5.6	36
48	Effects of single-agent and combination chemotherapy for gestational trophoblastic tumors on risks of second malignancy and early menopause. <i>Journal of Clinical Oncology</i> , 2015 , 33, 472-8	2.2	52
47	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	11	59
46	Anthropometric and hormonal risk factors for male breast cancer: male breast cancer pooling project results. <i>Journal of the National Cancer Institute</i> , 2014 , 106, djt465	9.7	92
45	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014 , 4, 4999	17.4	87
44	Genetic variation in mitotic regulatory pathway genes is associated with breast tumor grade. <i>Human Molecular Genetics</i> , 2014 , 23, 6034-46	5.6	11
43	Current knowledge and future research directions in treatment-related second primary malignancies. <i>European Journal of Cancer, Supplement</i> , 2014 , 12, 5-17	1.6	39
42	Genetic variation at CYP3A is associated with age at menarche and breast cancer risk: a case-control study. <i>Breast Cancer Research</i> , 2014 , 16, R51	8.3	12
41	MicroRNA related polymorphisms and breast cancer risk. <i>PLoS ONE</i> , 2014 , 9, e109973	3.7	37
40	Genetic predisposition to in situ and invasive lobular carcinoma of the breast. <i>PLoS Genetics</i> , 2014 , 10, e1004285	6	38
39	Risk of premature menopause after treatment for Hodgkin lymphoma. <i>Journal of the National Cancer Institute</i> , 2014 , 106,	9.7	37
38	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014 , 23, 6616-33	5.6	77
37	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2014 , 23, 6096-111	5.6	48
36	Radiologic features of breast cancer after mantle radiation therapy for Hodgkin disease: a study of 230 cases. <i>Radiology</i> , 2014 , 272, 73-8	20.5	10
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