

# John L. Hopper

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6495145/publications.pdf>

Version: 2024-02-01

531  
papers

34,733  
citations

6124

83  
h-index

7043

159  
g-index

547  
all docs

547  
docs citations

547  
times ranked

36264  
citing authors

#	ARTICLE	IF	CITATIONS
1	Blood DNA methylation score predicts breast cancer risk: applying OPERA in molecular, environmental, genetic and analytic epidemiology. <i>Molecular Oncology</i> , 2022, 16, 8-10.	2.1	3
2	Birthweight, gestational age and familial confounding in sex differences in infant mortality: a matched co-twin control study of Brazilian male-female twin pairs identified by population data linkage. <i>International Journal of Epidemiology</i> , 2022, 51, 1502-1510.	0.9	8
3	Alcohol and tobacco use and risk of multiple myeloma: A case-control study. <i>EJHaem</i> , 2022, 3, 109-120.	0.4	3
4	Risks of breast and ovarian cancer for women harboring pathogenic missense variants in BRCA1 and BRCA2 compared with those harboring protein truncating variants. <i>Genetics in Medicine</i> , 2022, 24, 119-129.	1.1	10
5	Cancer Risks Associated With <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variants. <i>Journal of Clinical Oncology</i> , 2022, 40, 1529-1541.	0.8	90
6	Rare germline copy number variants (CNVs) and breast cancer risk. <i>Communications Biology</i> , 2022, 5, 65.	2.0	6
7	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 349-362.	1.4	23
8	Common variants in breast cancer risk loci predispose to distinct tumor subtypes. <i>Breast Cancer Research</i> , 2022, 24, 2.	2.2	15
9	Oral Contraceptive Use in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers: Absolute Cancer Risks and Benefits. <i>Journal of the National Cancer Institute</i> , 2022, 114, 540-552.	3.0	7
10	Together Alone: Going Online during COVID-19 Is Changing Scientific Conferences. <i>Challenges</i> , 2022, 13, 7.	0.9	3
11	Association between very to moderate preterm births, lung function deficits, and COPD at age 53 years: analysis of a prospective cohort study. <i>Lancet Respiratory Medicine</i> , 2022, 10, 478-484.	5.2	42
12	Familial Aspects of Mammographic Density Measures Associated with Breast Cancer Risk. <i>Cancers</i> , 2022, 14, 1483.	1.7	6
13	Early life affects late-life health through determining DNA methylation across the lifespan: A twin study. <i>EBioMedicine</i> , 2022, 77, 103927.	2.7	15
14	Population-based estimates of age-specific cumulative risk of breast cancer for pathogenic variants in ATM. <i>Breast Cancer Research</i> , 2022, 24, 24.	2.2	3
15	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
16	Association of contralateral breast cancer risk with mammographic density defined at higher than conventional intensity thresholds. <i>International Journal of Cancer</i> , 2022, 151, 1304-1309.	2.3	3
17	Weight is More Informative than Body Mass Index for Predicting Postmenopausal Breast Cancer Risk: Prospective Family Study Cohort (ProF-SC). <i>Cancer Prevention Research</i> , 2022, 15, 185-191.	0.7	4
18	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

#	ARTICLE	IF	CITATIONS
19	Genetic Regulation of DNA Methylation Yields Novel Discoveries in GWAS of Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 1068-1076.	1.1	1
20	Within-sibship genome-wide association analyses decrease bias in estimates of direct genetic effects. <i>Nature Genetics</i> , 2022, 54, 581-592.	9.4	142
21	Reply to V. Fallet et al. <i>Journal of Clinical Oncology</i> , 2022, 40, 2509-2510.	0.8	3
22	Genetic Aspects of Mammographic Density Measures Associated with Breast Cancer Risk. <i>Cancers</i> , 2022, 14, 2767.	1.7	5
23	Does genetic predisposition modify the effect of lifestyle-related factors on DNA methylation?. <i>Epigenetics</i> , 2022, 17, 1838-1847.	1.3	2
24	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
25	The association of age at menarche and adult height with mammographic density in the International Consortium of Mammographic Density. <i>Breast Cancer Research</i> , 2022, 24, .	2.2	6
26	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
27	Novel mammogram-based measures improve breast cancer risk prediction beyond an established mammographic density measure. <i>International Journal of Cancer</i> , 2021, 148, 2193-2202.	2.3	18
28	Methylation marks of prenatal exposure to maternal smoking and risk of cancer in adulthood. <i>International Journal of Epidemiology</i> , 2021, 50, 105-115.	0.9	18
29	Comparing 5-Year and Lifetime Risks of Breast Cancer Using the Prospective Family Study Cohort. <i>Journal of the National Cancer Institute</i> , 2021, 113, 785-791.	3.0	13
30	DNA methylation and breast cancer risk: value of twin and family studies. , 2021, , 67-83.		1
31	Familial and Genetic Influences on the Common Pediatric Primary Pain Disorders: A Twin Family Study. <i>Children</i> , 2021, 8, 89.	0.6	5
32	Value of twin and family study designs for epigenetic research. , 2021, , 3-16.		0
33	CYP3A7*1C allele: linking premenopausal oestrogen and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , 2021, 124, 842-854.	2.9	5
34	Evaluating the utility of tumour mutational signatures for identifying hereditary colorectal cancer and polyposis syndrome carriers. <i>Gut</i> , 2021, 70, 2138-2149.	6.1	27
35	A streamlined model for use in clinical breast cancer risk assessment maintains predictive power and is further improved with inclusion of a polygenic risk score. <i>PLoS ONE</i> , 2021, 16, e0245375.	1.1	6
36	Sex differences in epigenetic profiles: The value of twin studies. , 2021, , 225-235.		0

#	ARTICLE	IF	CITATIONS
37	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
38	Genetic architectures of proximal and distal colorectal cancer are partly distinct. <i>Gut</i> , 2021, 70, 1325-1334.	6.1	44
39	Lifetime alcohol intake, drinking patterns over time and risk of stomach cancer: A pooled analysis of data from two prospective cohort studies. <i>International Journal of Cancer</i> , 2021, 148, 2759-2773.	2.3	7
40	Prospective Evaluation of the Addition of Polygenic Risk Scores to Breast Cancer Risk Models. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab021.	1.4	19
41	Germline and Tumor Sequencing as a Diagnostic Tool To Resolve Suspected Lynch Syndrome. <i>Journal of Molecular Diagnostics</i> , 2021, 23, 358-371.	1.2	12
42	VTRNA2-1: Genetic Variation, Heritable Methylation and Disease Association. <i>International Journal of Molecular Sciences</i> , 2021, 22, 2535.	1.8	15
43	Population-Based Estimates of the Age-Specific Cumulative Risk of Breast Cancer for Pathogenic Variants in CHEK2: Findings from the Australian Breast Cancer Family Registry. <i>Cancers</i> , 2021, 13, 1378.	1.7	5
44	Age dependency of the polygenic risk score for colorectal cancer. <i>American Journal of Human Genetics</i> , 2021, 108, 525-526.	2.6	12
45	Assessment of a Polygenic Risk Score for Colorectal Cancer to Predict Risk of Lynch Syndrome Colorectal Cancer. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab022.	1.4	15
46	Epigenetic Drift Association with Cancer Risk and Survival, and Modification by Sex. <i>Cancers</i> , 2021, 13, 1881.	1.7	9
47	Uptake of hysterectomy and bilateral salpingo-oophorectomy in carriers of pathogenic mismatch repair variants: a Prospective Lynch Syndrome Database report. <i>European Journal of Cancer</i> , 2021, 148, 124-133.	1.3	11
48	Evaluation of the association of heterozygous germline variants in NTHL1 with breast cancer predisposition: an international multi-center study of 47,180 subjects. <i>Npj Breast Cancer</i> , 2021, 7, 52.	2.3	7
49	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
50	Prediagnosis alcohol intake and metachronous cancer risk in cancer survivors: A prospective cohort study. <i>International Journal of Cancer</i> , 2021, 149, 827-838.	2.3	2
51	DNA Methylation Signatures and the Contribution of Age-Associated Methylomic Drift to Carcinogenesis in Early-Onset Colorectal Cancer. <i>Cancers</i> , 2021, 13, 2589.	1.7	18
52	Association between Smoking and Molecular Subtypes of Colorectal Cancer. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab056.	1.4	8
53	RE: Chemopreventive Agents to Reduce Mammographic Breast Density in Premenopausal Women: A Systematic Review of Clinical Trials. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab051.	1.4	1
54	The predictive ability of the 313 variant-based polygenic risk score for contralateral breast cancer risk prediction in women of European ancestry with a heterozygous BRCA1 or BRCA2 pathogenic variant. <i>Genetics in Medicine</i> , 2021, 23, 1726-1737.	1.1	16

#	ARTICLE	IF	CITATIONS
55	No Difference in Penetrance between Truncating and Missense/Aberrant Splicing Pathogenic Variants in MLH1 and MSH2: A Prospective Lynch Syndrome Database Study. <i>Journal of Clinical Medicine</i> , 2021, 10, 2856.	1.0	11
56	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
57	Tetranucleotide and Low Microsatellite Instability Are Inversely Associated with the CpG Island Methylator Phenotype in Colorectal Cancer. <i>Cancers</i> , 2021, 13, 3529.	1.7	3
58	Oral contraceptive use and ovarian cancer risk for BRCA1/2 mutation carriers: an international cohort study. <i>American Journal of Obstetrics and Gynecology</i> , 2021, 225, 51.e1-51.e17.	0.7	34
59	Surrounding Greenness and Biological Aging Based on DNA Methylation: A Twin and Family Study in Australia. <i>Environmental Health Perspectives</i> , 2021, 129, 87007.	2.8	14
60	Residential surrounding greenness and DNA methylation: an epigenome-wide association study. <i>ISEE Conference Abstracts</i> , 2021, 2021, .	0.0	0
61	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
62	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
63	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	13.7	183
64	Surrounding greenness is associated with slower biological ageing based on epigenetics. <i>International Journal of Epidemiology</i> , 2021, 50, .	0.9	0
65	Ambient temperature and genome-wide DNA methylation: A twin and family study in Australia. <i>Environmental Pollution</i> , 2021, 285, 117700.	3.7	9
66	Towards risk-stratified population breast cancer screening: more than mammographic density. <i>Medical Journal of Australia</i> , 2021, 215, 350-351.	0.8	2
67	Inference on Causation from Examining Changes in Regression coefficients and Innovative Statistical Analyses (ICE CRISTAL). <i>International Journal of Epidemiology</i> , 2021, 50, .	0.9	0
68	Discriminating between risk discriminators: OPERA, AUC, and polygenic variance. <i>International Journal of Epidemiology</i> , 2021, 50, .	0.9	0
69	ICE FALCON: a causation assessment method analogous to, but more powerful than, Mendelian Randomisation. <i>International Journal of Epidemiology</i> , 2021, 50, .	0.9	0
70	Residential surrounding greenness and DNA methylation: An epigenome-wide association study. <i>Environment International</i> , 2021, 154, 106556.	4.8	23
71	Smoking Methylation Marks for Prediction of Urothelial Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 2197-2206.	1.1	4
72	Novel approach to estimating sex differences unconfounded by familial factors from studying male-female twin pairs. <i>International Journal of Epidemiology</i> , 2021, 50, .	0.9	1

#	ARTICLE	IF	CITATIONS
73	32Do the risks of Lynch syndrome-related cancers depend on the parent-of-origin of the mutation?. International Journal of Epidemiology, 2021, 50, .	0.9	0
74	Ability of known colorectal cancer susceptibility SNPs to predict colorectal cancer risk: A cohort study within the UK Biobank. PLoS ONE, 2021, 16, e0251469.	1.1	5
75	Smoking, alcohol consumption, body fatness, and risk of myelodysplastic syndromes: A prospective study. Leukemia Research, 2021, 109, 106593.	0.4	1
76	Educational attainment of same-sex and opposite-sex dizygotic twins: An individual-level pooled study of 19 twin cohorts. Hormones and Behavior, 2021, 136, 105054.	1.0	1
77	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
78	Biological Aging Measures Based on Blood DNA Methylation and Risk of Cancer: A Prospective Study. JNCI Cancer Spectrum, 2021, 5, pkaa109.	1.4	40
79	A Combined Proteomics and Mendelian Randomization Approach to Investigate the Effects of Aspirin-Targeted Proteins on Colorectal Cancer. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 564-575.	1.1	10
80	Breast Cancer Chemoprevention: Use and Views of Australian Women and Their Clinicians. Cancer Prevention Research, 2021, 14, 131-144.	0.7	6
81	Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. Scientific Reports, 2021, 11, 19787.	1.6	2
82	Prospective Evaluation over 15 Years of Six Breast Cancer Risk Models. Cancers, 2021, 13, 5194.	1.7	7
83	Repeatability of methylation measures using a QIaseq targeted methyl panel and comparison with the Illumina HumanMethylation450 assay. BMC Research Notes, 2021, 14, 394.	0.6	2
84	Motivators of Inappropriate Ovarian Cancer Screening: A Survey of Women and Their Clinicians. JNCI Cancer Spectrum, 2021, 5, pkaa110.	1.4	4
85	Salicylic Acid and Risk of Colorectal Cancer: A Two-Sample Mendelian Randomization Study. Nutrients, 2021, 13, 4164.	1.7	3
86	Mammographic texture features associated with contralateral breast cancer in the WECARE Study. Npj Breast Cancer, 2021, 7, 146.	2.3	1
87	Association of chronic musculoskeletal pain with mortality among UK adults: A population-based cohort study with mediation analysis. EClinicalMedicine, 2021, 42, 101202.	3.2	6
88	Association of FOXO3 Blood DNA Methylation with Cancer Risk, Cancer Survival, and Mortality. Cells, 2021, 10, 3384.	1.8	6
89	Recreational Physical Activity and Outcomes After Breast Cancer in Women at High Familial Risk. JNCI Cancer Spectrum, 2021, 5, pkab090.	1.4	1
90	Population-based estimates of breast cancer risk for carriers of pathogenic variants identified by gene-panel testing. Npj Breast Cancer, 2021, 7, 153.	2.3	10

#	ARTICLE	IF	CITATIONS
91	Association of Genomic Domains in <i>BRCA1</i> and <i>BRCA2</i> with Prostate Cancer Risk and Aggressiveness. <i>Cancer Research</i> , 2020, 80, 624-638.	0.4	39
92	Considerations When Using Breast Cancer Risk Models for Women with Negative <i>BRCA1/BRCA2</i> Mutation Results. <i>Journal of the National Cancer Institute</i> , 2020, 112, 418-422.	3.0	1
93	Recreational Physical Activity Is Associated with Reduced Breast Cancer Risk in Adult Women at High Risk for Breast Cancer: A Cohort Study of Women Selected for Familial and Genetic Risk. <i>Cancer Research</i> , 2020, 80, 116-125.	0.4	37
94	Interval breast cancer risk associations with breast density, family history and breast tissue aging. <i>International Journal of Cancer</i> , 2020, 147, 375-382.	2.3	22
95	Early birth is a key factor in educational disadvantage of twins: A data linkage study. <i>Acta Paediatrica, International Journal of Paediatrics</i> , 2020, 109, 534-540.	0.7	3
96	Cancer risks by gene, age, and gender in 6350 carriers of pathogenic mismatch repair variants: findings from the Prospective Lynch Syndrome Database. <i>Genetics in Medicine</i> , 2020, 22, 15-25.	1.1	365
97	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
98	Overall lack of replication of associations between dietary intake of folate and vitamin B-12 and DNA methylation in peripheral blood. <i>American Journal of Clinical Nutrition</i> , 2020, 111, 228-230.	2.2	6
99	Cumulative Burden of Colorectal Cancer-Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. <i>Gastroenterology</i> , 2020, 158, 1274-1286.e12.	0.6	110
100	Cancer Risks Associated With Germline <i>PALB2</i> Pathogenic Variants: An International Study of 524 Families. <i>Journal of Clinical Oncology</i> , 2020, 38, 674-685.	0.8	270
101	Economic Evaluation of Population-Based <i>BRCA1/BRCA2</i> Mutation Testing across Multiple Countries and Health Systems. <i>Cancers</i> , 2020, 12, 1929.	1.7	49
102	Are the Relationships of Lean Mass and Fat Mass With Bone Microarchitecture Causal or Due to Familial Confounders? A Novel Study of Adult Female Twin Pairs. <i>JBMR Plus</i> , 2020, 4, e10386.	1.3	6
103	Inference about causation from examination of familial confounding (ICE FALCON): a model for assessing causation analogous to Mendelian randomization. <i>International Journal of Epidemiology</i> , 2020, 49, 1259-1269.	0.9	26
104	Prognostic value of metabolic tumor volume and total lesion glycolysis in breast cancer: a meta-analysis. <i>Nuclear Medicine Communications</i> , 2020, 41, 824-829.	0.5	9
105	Stochastic Epigenetic Mutations Are Associated with Risk of Breast Cancer, Lung Cancer, and Mature B-cell Neoplasms. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2026-2037.	1.1	18
106	Genetic and environmental variation in educational attainment: an individual-based analysis of 28 twin cohorts. <i>Scientific Reports</i> , 2020, 10, 12681.	1.6	59
107	Avoiding dynastic, assortative mating, and population stratification biases in Mendelian randomization through within-family analyses. <i>Nature Communications</i> , 2020, 11, 3519.	5.8	213
108	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.	2.6	39

#	ARTICLE	IF	CITATIONS
109	Circulating bilirubin levels and risk of colorectal cancer: serological and Mendelian randomization analyses. <i>BMC Medicine</i> , 2020, 18, 229.	2.3	28
110	Contrasting painless and painful phenotypes of pediatric restless legs syndrome: a twin family study. <i>Sleep Medicine</i> , 2020, 75, 361-367.	0.8	7
111	Genetic and environmental causes of variation in epigenetic aging across the lifespan. <i>Clinical Epigenetics</i> , 2020, 12, 158.	1.8	33
112	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
113	Genetic and environmental influences on human height from infancy through adulthood at different levels of parental education. <i>Scientific Reports</i> , 2020, 10, 7974.	1.6	17
114	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
115	Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 860-870.	1.1	26
116	Characterization of the Cancer Spectrum in Men With Germline <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variants. <i>JAMA Oncology</i> , 2020, 6, 1218.	3.4	48
117	Do the risks of Lynch syndrome-related cancers depend on the parent of origin of the mutation?. <i>Familial Cancer</i> , 2020, 19, 215-222.	0.9	1
118	Going Beyond Conventional Mammographic Density to Discover Novel Mammogram-Based Predictors of Breast Cancer Risk. <i>Journal of Clinical Medicine</i> , 2020, 9, 627.	1.0	23
119	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	0.6	32
120	Mismatch repair gene pathogenic germline variants in a population-based cohort of breast cancer. <i>Familial Cancer</i> , 2020, 19, 197-202.	0.9	6
121	Alcohol Consumption, Cigarette Smoking, and Risk of Breast Cancer for <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers: Results from The <i>BRCA1</i> and <i>BRCA2</i> Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 368-378.	1.1	24
122	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312.	5.8	30
123	Risk-reducing salpingo-oophorectomy, natural menopause, and breast cancer risk: an international prospective cohort of <i>BRCA1</i> and <i>BRCA2</i> mutation carriers. <i>Breast Cancer Research</i> , 2020, 22, 8.	2.2	41
124	A New Comprehensive Colorectal Cancer Risk Prediction Model Incorporating Family History, Personal Characteristics, and Environmental Factors. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 549-557.	1.1	25
125	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
126	Physical activity and risks of breast and colorectal cancer: a Mendelian randomisation analysis. <i>Nature Communications</i> , 2020, 11, 597.	5.8	193



#	ARTICLE	IF	CITATIONS
127	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 146-157.	3.0	129
128	Inference about causation between body mass index and DNA methylation in blood from a twin family study. <i>International Journal of Obesity</i> , 2019, 43, 243-252.	1.6	48
129	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
130	DNA methylation-based biological age, genome-wide average DNA methylation, and conventional breast cancer risk factors. <i>Scientific Reports</i> , 2019, 9, 15055.	1.6	18
131	Accuracy of Risk Estimates from the iPrevent Breast Cancer Risk Assessment and Management Tool. <i>JNCI Cancer Spectrum</i> , 2019, 3, pkz066.	1.4	8
132	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , 2019, 9, 12524.	1.6	5
133	A Cost-effectiveness Analysis of Multigene Testing for All Patients With Breast Cancer. <i>JAMA Oncology</i> , 2019, 5, 1718.	3.4	91
134	Type 2 diabetes mellitus, blood cholesterol, triglyceride and colorectal cancer risk in Lynch syndrome. <i>British Journal of Cancer</i> , 2019, 121, 869-876.	2.9	10
135	Body size and dietary risk factors for aggressive prostate cancer: a case-control study. <i>Cancer Causes and Control</i> , 2019, 30, 1301-1312.	0.8	2
136	Bivariate mixture models for the joint distribution of repeated serum ferritin and transferrin saturation measured 12 years apart in a cohort of healthy middle-aged Australians. <i>PLoS ONE</i> , 2019, 14, e0214196.	1.1	0
137	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88
138	Genome-wide association study of peripheral blood DNA methylation and conventional mammographic density measures. <i>International Journal of Cancer</i> , 2019, 145, 1768-1773.	2.3	17
139	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
140	Ability of known susceptibility SNPs to predict colorectal cancer risk for persons with and without a family history. <i>Familial Cancer</i> , 2019, 18, 389-397.	0.9	23
141	Mendelian randomisation study of height and body mass index as modifiers of ovarian cancer risk in 22,588 BRCA1 and BRCA2 mutation carriers. <i>British Journal of Cancer</i> , 2019, 121, 180-192.	2.9	19
142	Performance of BCRAT in high-risk patients with breast cancer – Authors' reply. <i>Lancet Oncology</i> , The, 2019, 20, e286.	5.1	3
143	Association analyses identify 31 new risk loci for colorectal cancer susceptibility. <i>Nature Communications</i> , 2019, 10, 2154.	5.8	172
144	Regular use of aspirin and other non-steroidal anti-inflammatory drugs and breast cancer risk for women at familial or genetic risk: a cohort study. <i>Breast Cancer Research</i> , 2019, 21, 52.	2.2	44

#	ARTICLE	IF	CITATIONS
145	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
146	Occupational exposure to solvents and lung function decline: A population based study. <i>Thorax</i> , 2019, 74, 650-658.	2.7	21
147	Parental Education and Genetics of BMI from Infancy to Old Age: A Pooled Analysis of 29 Twin Cohorts. <i>Obesity</i> , 2019, 27, 855-865.	1.5	27
148	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
149	Research Note: Twin studies and their value for physiotherapy research. <i>Journal of Physiotherapy</i> , 2019, 65, 58-60.	0.7	3
150	Lifetime alcohol intake and pancreatic cancer incidence and survival: findings from the Melbourne Collaborative Cohort Study. <i>Cancer Causes and Control</i> , 2019, 30, 323-331.	0.8	7
151	Benign breast disease increases breast cancer risk independent of underlying familial risk profile: Findings from a Prospective Family Study Cohort. <i>International Journal of Cancer</i> , 2019, 145, 370-379.	2.3	9
152	Mortality after breast cancer as a function of time since diagnosis by estrogen receptor status and age at diagnosis. <i>International Journal of Cancer</i> , 2019, 145, 3207-3217.	2.3	14
153	10-year performance of four models of breast cancer risk: a validation study. <i>Lancet Oncology</i> , The, 2019, 20, 504-517.	5.1	116
154	Measurement challenge: protocol for international case-control comparison of mammographic measures that predict breast cancer risk. <i>BMJ Open</i> , 2019, 9, e031041.	0.8	14
155	Prediction and clinical utility of a contralateral breast cancer risk model. <i>Breast Cancer Research</i> , 2019, 21, 144.	2.2	24
156	Alcohol consumption, cigarette smoking, and familial breast cancer risk: findings from the Prospective Family Study Cohort (ProF-SC). <i>Breast Cancer Research</i> , 2019, 21, 128.	2.2	27
157	Twins Research Australia: A New Paradigm for Driving Twin Research. <i>Twin Research and Human Genetics</i> , 2019, 22, 438-445.	0.3	17
158	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019, 104, 21-34.	2.6	711
159	Debunking a myopic view of nature versus nurture. <i>Australasian journal of optometry</i> , The, 2019, 102, 1-2.	0.6	9
160	Height and Body Mass Index as Modifiers of Breast Cancer Risk in <i>BRCA1</i> Mutation Carriers: A Mendelian Randomization Study. <i>Journal of the National Cancer Institute</i> , 2019, 111, 350-364.	3.0	30
161	Discovery of common and rare genetic risk variants for colorectal cancer. <i>Nature Genetics</i> , 2019, 51, 76-87.	9.4	377
162	Risk-Reducing Oophorectomy and Breast Cancer Risk Across the Spectrum of Familial Risk. <i>Journal of the National Cancer Institute</i> , 2019, 111, 331-334.	3.0	31

#	ARTICLE	IF	CITATIONS
163	Assessing the ProMCol classifier as a prognostic marker for non-metastatic colorectal cancer within the Melbourne Collaborative Cohort Study. <i>Gut</i> , 2019, 68, 761-762.	6.1	2
164	The <i>BRCA2</i> c.68-7T>A variant is not pathogenic: A model for clinical calibration of spliceogenicity. <i>Human Mutation</i> , 2018, 39, 729-741.	1.1	19
165	Cohort Profile: The Colon Cancer Family Registry Cohort (CCFRC). <i>International Journal of Epidemiology</i> , 2018, 47, 387-388i.	0.9	40
166	Heritable DNA methylation marks associated with susceptibility to breast cancer. <i>Nature Communications</i> , 2018, 9, 867.	5.8	76
167	Determining Risk of Colorectal Cancer and Starting Age of Screening Based on Lifestyle, Environmental, and Genetic Factors. <i>Gastroenterology</i> , 2018, 154, 2152-2164.e19.	0.6	226
168	Birth size and gestational age in opposite-sex twins as compared to same-sex twins: An individual-based pooled analysis of 21 cohorts. <i>Scientific Reports</i> , 2018, 8, 6300.	1.6	21
169	Associations between birth size and later height from infancy through adulthood: An individual based pooled analysis of 28 twin cohorts participating in the CODAtwins project. <i>Early Human Development</i> , 2018, 120, 53-60.	0.8	20
170	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.	1.3	62
171	Childhood measles contributes to post-bronchodilator airflow obstruction in middle-aged adults: A cohort study. <i>Respirology</i> , 2018, 23, 780-787.	1.3	5
172	An open-source, integrated pedigree data management and visualization tool for genetic epidemiology. <i>International Journal of Epidemiology</i> , 2018, 47, 1034-1039.	0.9	5
173	Dietary intake of nutrients involved in one-carbon metabolism and risk of urothelial cell carcinoma: A prospective cohort study. <i>International Journal of Cancer</i> , 2018, 143, 298-306.	2.3	12
174	Cost-effectiveness of Population-Based BRCA1, BRCA2, RAD51C, RAD51D, BRIP1, PALB2 Mutation Testing in Unselected General Population Women. <i>Journal of the National Cancer Institute</i> , 2018, 110, 714-725.	3.0	138
175	Twin studies for the prognosis, prevention and treatment of musculoskeletal conditions. <i>Brazilian Journal of Physical Therapy</i> , 2018, 22, 184-189.	1.1	9
176	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.	0.9	88
177	Is RNASEL:p.Glu265* a modifier of early-onset breast cancer risk for carriers of high-risk mutations?. <i>BMC Cancer</i> , 2018, 18, 165.	1.1	6
178	A Mixed-Effects Model for Powerful Association Tests in Integrative Functional Genomics. <i>American Journal of Human Genetics</i> , 2018, 102, 904-919.	2.6	30
179	Somatic mutations of the coding microsatellites within the beta-2-microglobulin gene in mismatch repair-deficient colorectal cancers and adenomas. <i>Familial Cancer</i> , 2018, 17, 91-100.	0.9	21
180	Lifetime alcohol intake and risk of non-Hodgkin lymphoma: Findings from the Melbourne Collaborative Cohort Study. <i>International Journal of Cancer</i> , 2018, 142, 919-926.	2.3	6

#	ARTICLE	IF	CITATIONS
181	Breast Cancer Risk Associations with Digital Mammographic Density by Pixel Brightness Threshold and Mammographic System. <i>Radiology</i> , 2018, 286, 433-442.	3.6	29
182	Association between mammographic density and tumor marker-defined breast cancer subtypes: a case-control study. <i>European Journal of Cancer Prevention</i> , 2018, 27, 239-247.	0.6	13
183	Genome-wide association study and meta-analysis in Northern European populations replicate multiple colorectal cancer risk loci. <i>International Journal of Cancer</i> , 2018, 142, 540-546.	2.3	26
184	Association of DNA Methylation-Based Biological Age With Health Risk Factors and Overall and Cause-Specific Mortality. <i>American Journal of Epidemiology</i> , 2018, 187, 529-538.	1.6	106
185	Associations of alcohol intake, smoking, physical activity and obesity with survival following colorectal cancer diagnosis by stage, anatomic site and tumor molecular subtype. <i>International Journal of Cancer</i> , 2018, 142, 238-250.	2.3	83
186	DNA methylation-based biological aging and cancer risk and survival: Pooled analysis of seven prospective studies. <i>International Journal of Cancer</i> , 2018, 142, 1611-1619.	2.3	153
187	Menopause-Related Appendicular Bone Loss is Mainly Cortical and Results in Increased Cortical Porosity. <i>Journal of Bone and Mineral Research</i> , 2018, 33, 598-605.	3.1	37
188	Risk of colorectal cancer for carriers of a germ-line mutation in POLE or POLD1. <i>Genetics in Medicine</i> , 2018, 20, 890-895.	1.1	49
189	Cancer Risks for PMS2-Associated Lynch Syndrome. <i>Journal of Clinical Oncology</i> , 2018, 36, 2961-2968.	0.8	147
190	Cirrus: An Automated Mammography-Based Measure of Breast Cancer Risk Based on Textural Features. <i>JNCI Cancer Spectrum</i> , 2018, 2, pky057.	1.4	24
191	Genome-wide DNA methylation assessment of BRCA1-like early-onset breast cancer: Data from the Australian Breast Cancer Family Registry. <i>Experimental and Molecular Pathology</i> , 2018, 105, 404-410.	0.9	26
192	Age-specific breast cancer risk by body mass index and familial risk: prospective family study cohort (ProF-SC). <i>Breast Cancer Research</i> , 2018, 20, 132.	2.2	51
193	Predicting interval and screen-detected breast cancers from mammographic density defined by different brightness thresholds. <i>Breast Cancer Research</i> , 2018, 20, 152.	2.2	24
194	Validation of a genetic risk score for Arkansas women of color. <i>PLoS ONE</i> , 2018, 13, e0204834.	1.1	12
195	Oral Contraceptive Use and Breast Cancer Risk: Retrospective and Prospective Analyses From a BRCA1 and BRCA2 Mutation Carrier Cohort Study. <i>JNCI Cancer Spectrum</i> , 2018, 2, pky023.	1.4	33
196	InforMD: a new initiative to raise public awareness about breast density. <i>Ecancermedalscience</i> , 2018, 12, 807.	0.6	4
197	BRCA1/2-negative, high-risk breast cancers (BRCAX) for Asian women: genetic susceptibility loci and their potential impacts. <i>Scientific Reports</i> , 2018, 8, 15263.	1.6	25
198	Genome-wide average DNA methylation is determined in utero. <i>International Journal of Epidemiology</i> , 2018, 47, 908-916.	0.9	38

#	ARTICLE	IF	CITATIONS
199	Genetic susceptibility markers for a breast-colorectal cancer phenotype: Exploratory results from genome-wide association studies. <i>PLoS ONE</i> , 2018, 13, e0196245.	1.1	9
200	Genetic and environmental factors affecting birth size variation: a pooled individual-based analysis of secular trends and global geographical differences using 26 twin cohorts. <i>International Journal of Epidemiology</i> , 2018, 47, 1195-1206.	0.9	19
201	Mendelian randomisation study of age at menarche and age at menopause and the risk of colorectal cancer. <i>British Journal of Cancer</i> , 2018, 118, 1639-1647.	2.9	16
202	DNA Methylation-Based Measures of Biological Aging. , 2018, , 39-64.		16
203	Association of current and former smoking with body mass index: A study of smoking discordant twin pairs from 21 twin cohorts. <i>PLoS ONE</i> , 2018, 13, e0200140.	1.1	57
204	Sunscreen Use and Melanoma Risk Among Young Australian Adults. <i>JAMA Dermatology</i> , 2018, 154, 1001.	2.0	40
205	Causal effect of smoking on DNA methylation in peripheral blood: a twin and family study. <i>Clinical Epigenetics</i> , 2018, 10, 18.	1.8	95
206	Epigenetic supersimilarity of monozygotic twin pairs. <i>Genome Biology</i> , 2018, 19, 2.	3.8	89
207	Family history-based colorectal cancer screening in Australia: A modelling study of the costs, benefits, and harms of different participation scenarios. <i>PLoS Medicine</i> , 2018, 15, e1002630.	3.9	6
208	Mammographic Density and Circulating Sex Hormones: a Cross-Sectional Study in Postmenopausal Korean Women. <i>Hormones and Cancer</i> , 2018, 9, 383-390.	4.9	2
209	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018, 9, 3166.	5.8	178
210	Assessing the Incremental Contribution of Common Genomic Variants to Melanoma Risk Prediction in Two Population-Based Studies. <i>Journal of Investigative Dermatology</i> , 2018, 138, 2617-2624.	0.3	52
211	Association between birth weight and educational attainment: an individual-based pooled analysis of nine twin cohorts. <i>Journal of Epidemiology and Community Health</i> , 2018, 72, 832-837.	2.0	5
212	Physical activity and the risk of colorectal cancer in Lynch syndrome. <i>International Journal of Cancer</i> , 2018, 143, 2250-2260.	2.3	23
213	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.	9.4	184
214	The iPrevent Online Breast Cancer Risk Assessment and Risk Management Tool: Usability and Acceptability Testing. <i>JMIR Formative Research</i> , 2018, 2, e24.	0.7	10
215	Cohort Profile: The Tasmanian Longitudinal Health STUDY (TAHS). <i>International Journal of Epidemiology</i> , 2017, 46, dyw028.	0.9	26
216	The Ark: a customizable web-based data management tool for health and medical research. <i>Bioinformatics</i> , 2017, 33, 624-626.	1.8	6

#	ARTICLE	IF	CITATIONS
217	Mammographic density defined by higher than conventional brightness thresholds better predicts breast cancer risk. <i>International Journal of Epidemiology</i> , 2017, 46, dyw212.	0.9	24
218	Breast cancer risk prediction using a polygenic risk score in the familial setting: a prospective study from the Breast Cancer Family Registry and kConFab. <i>Genetics in Medicine</i> , 2017, 19, 30-35.	1.1	53
219	Tumor testing to identify lynch syndrome in two Australian colorectal cancer cohorts. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2017, 32, 427-438.	1.4	47
220	Longitudinal Study of Mammographic Density Measures That Predict Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 651-660.	1.1	36
221	Genome-Wide Measures of Peripheral Blood Dna Methylation and Prostate Cancer Risk in a Prospective Nested Case-Control Study. <i>Prostate</i> , 2017, 77, 471-478.	1.2	31
222	<i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017, 77, 2789-2799.	0.4	75
223	Common Pediatric Pain Disorders and Their Clinical Associations. <i>Clinical Journal of Pain</i> , 2017, 33, 1131-1140.	0.8	13
224	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
225	Genetics for population and public health. <i>International Journal of Epidemiology</i> , 2017, 46, 8-11.	0.9	12
226	Risks of Breast, Ovarian, and Contralateral Breast Cancer for <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>JAMA - Journal of the American Medical Association</i> , 2017, 317, 2402.	3.8	1,898
227	Mendelian randomisation implicates hyperlipidaemia as a risk factor for colorectal cancer. <i>International Journal of Cancer</i> , 2017, 140, 2701-2708.	2.3	76
228	Association between birthweight and later body mass index: an individual-based pooled analysis of 27 twin cohorts participating in the CODATwins project. <i>International Journal of Epidemiology</i> , 2017, 46, 1488-1498.	0.9	22
229	Ejaculatory frequency and the risk of aggressive prostate cancer: Findings from a case-control study. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2017, 35, 530.e7-530.e13.	0.8	13
230	Dependence of cancer risk from environmental exposures on underlying genetic susceptibility: an illustration with polycyclic aromatic hydrocarbons and breast cancer. <i>British Journal of Cancer</i> , 2017, 116, 1229-1233.	2.9	54
231	The interaction between farming/rural environment and TLR2, TLR4, TLR6 and CD14 genetic polymorphisms in relation to early- and late-onset asthma. <i>Scientific Reports</i> , 2017, 7, 43681.	1.6	27
232	Genetic and Environmental Factors in Invasive Cervical Cancer: Design and Methods of a Classical Twin Study. <i>Twin Research and Human Genetics</i> , 2017, 20, 10-18.	0.3	11
233	Education in Twins and Their Parents Across Birth Cohorts Over 100 years: An Individual-Level Pooled Analysis of 42-Twin Cohorts. <i>Twin Research and Human Genetics</i> , 2017, 20, 395-405.	0.3	8
234	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099

#	ARTICLE	IF	CITATIONS
235	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
236	Pro-inflammatory fatty acid profile and colorectal cancer risk: A Mendelian randomisation analysis. <i>European Journal of Cancer</i> , 2017, 84, 228-238.	1.3	81
237	Twin birth changes DNA methylation of subsequent siblings. <i>Scientific Reports</i> , 2017, 7, 8463.	1.6	8
238	Bronchial hyperresponsiveness and obesity in middle age: insights from an Australian cohort. <i>European Respiratory Journal</i> , 2017, 50, 1602181.	3.1	20
239	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
240	Germline variation in ADAMTSL1 is associated with prognosis following breast cancer treatment in young women. <i>Nature Communications</i> , 2017, 8, 1632.	5.8	18
241	Causes of blood methylomic variation for middle-aged women measured by the HumanMethylation450 array. <i>Epigenetics</i> , 2017, 12, 973-981.	1.3	14
242	Testing for Gene-Environment Interactions Using a Prospective Family Cohort Design: Body Mass Index in Early and Later Adulthood and Risk of Breast Cancer. <i>American Journal of Epidemiology</i> , 2017, 185, 487-500.	1.6	5
243	Cost-effectiveness of population based BRCA testing with varying Ashkenazi Jewish ancestry. <i>American Journal of Obstetrics and Gynecology</i> , 2017, 217, 578.e1-578.e12.	0.7	63
244	Occupational exposure to pesticides are associated with fixed airflow obstruction in middle-age. <i>Thorax</i> , 2017, 72, 990-997.	2.7	32
245	Increased genomic burden of germline copy number variants is associated with early onset breast cancer: Australian breast cancer family registry. <i>Breast Cancer Research</i> , 2017, 19, 30.	2.2	14
246	Does the sex of one's co-twin affect height and BMI in adulthood? A study of dizygotic adult twins from 31 cohorts. <i>Biology of Sex Differences</i> , 2017, 8, 14.	1.8	8
247	Preterm birth and low birth weight continue to increase the risk of asthma from age 7 to 43. <i>Journal of Asthma</i> , 2017, 54, 616-623.	0.9	31
248	Alcohol Consumption and the Risk of Colorectal Cancer for Mismatch Repair Gene Mutation Carriers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 366-375.	1.1	37
249	<i>HFE</i> p.C282Y homozygosity predisposes to rapid serum ferritin rise after menopause: A genotype-stratified cohort study of hemochromatosis in Australian women. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2017, 32, 797-802.	1.4	16
250	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017, 19, 599-603.	1.1	67
251	Prevalence and Penetrance of Major Genes and Polygenes for Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 404-412.	1.1	341
252	Germline miRNA DNA variants and the risk of colorectal cancer by subtype. <i>Genes Chromosomes and Cancer</i> , 2017, 56, 177-184.	1.5	7

#	ARTICLE	IF	CITATIONS
253	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017, 46, 1814-1822.	0.9	45
254	Enrichment of colorectal cancer associations in functional regions: Insight for using epigenomics data in the analysis of whole genome sequence-imputed GWAS data. <i>PLoS ONE</i> , 2017, 12, e0186518.	1.1	8
255	Comparison of the association of mammographic density and clinical factors with ductal carcinoma in situ versus invasive ductal breast cancer in Korean women. <i>BMC Cancer</i> , 2017, 17, 821.	1.1	5
256	Mammographic density and risk of breast cancer by tumor characteristics: a case-control study. <i>BMC Cancer</i> , 2017, 17, 859.	1.1	5
257	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. <i>Breast Cancer Research</i> , 2017, 19, 119.	2.2	43
258	Reproductive factors as risk modifiers of breast cancer in <i>BRCA</i> mutation carriers and high-risk non-carriers. <i>Oncotarget</i> , 2017, 8, 102110-102118.	0.8	23
259	Mammographic density and ageing: A collaborative pooled analysis of cross-sectional data from 22 countries worldwide. <i>PLoS Medicine</i> , 2017, 14, e1002335.	3.9	108
260	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
261	<i>PHIP</i> - a novel candidate breast cancer susceptibility locus on 6q14.1. <i>Oncotarget</i> , 2017, 8, 102769-102782.	0.8	9
262	Current asthma contributes as much as smoking to chronic bronchitis in middle age: a prospective population-based study. <i>International Journal of COPD</i> , 2016, Volume 11, 1911-1920.	0.9	10
263	Genetic and environmental influences on adult human height across birth cohorts from 1886 to 1994. <i>ELife</i> , 2016, 5, .	2.8	42
264	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
265	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
266	ABRAXAS (FAM175A) and Breast Cancer Susceptibility: No Evidence of Association in the Breast Cancer Family Registry. <i>PLoS ONE</i> , 2016, 11, e0156820.	1.1	5
267	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. <i>PLoS ONE</i> , 2016, 11, e0160316.	1.1	12
268	Risk factors for metachronous colorectal cancer following a primary colorectal cancer: A prospective cohort study. <i>International Journal of Cancer</i> , 2016, 139, 1081-1090.	2.3	32
269	Mendelian randomisation analysis strongly implicates adiposity with risk of developing colorectal cancer. <i>British Journal of Cancer</i> , 2016, 115, 266-272.	2.9	57
270	Association between selected dietary scores and the risk of urothelial cell carcinoma: A prospective cohort study. <i>International Journal of Cancer</i> , 2016, 139, 1251-1260.	2.3	47



#	ARTICLE	IF	CITATIONS
271	Cholecystectomy and the risk of colorectal cancer by tumor mismatch repair deficiency status. <i>International Journal of Colorectal Disease</i> , 2016, 31, 1451-1457.	1.0	6
272	Clinical and functional differences between early-onset and late-onset adult asthma: a population-based Tasmanian Longitudinal Health Study. <i>Thorax</i> , 2016, 71, 981-987.	2.7	51
273	<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.	1.5	174
274	Transitioning to routine breast cancer risk assessment and management in primary care: what can we learn from cardiovascular disease?. <i>Australian Journal of Primary Health</i> , 2016, 22, 255.	0.4	16
275	Mammographic density and risk of breast cancer by mode of detection and tumor size: a case-control study. <i>Breast Cancer Research</i> , 2016, 18, 63.	2.2	30
276	The Brazilian Twin Registry. <i>Twin Research and Human Genetics</i> , 2016, 19, 687-691.	0.3	12
277	Protective and Harmful Effects of Physical Activity for Low Back Pain: A Protocol for the AUstralian Twin BACK Pain (AUTBACK) Feasibility Study. <i>Twin Research and Human Genetics</i> , 2016, 19, 502-509.	0.3	7
278	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.	2.2	31
279	Efficacy of a Sleep Quality Intervention in People With Low Back Pain: Protocol for a Feasibility Randomized Co-Twin Controlled Trial. <i>Twin Research and Human Genetics</i> , 2016, 19, 492-501.	0.3	16
280	Twin's Birth-Order Differences in Height and Body Mass Index From Birth to Old Age: A Pooled Study of 26 Twin Cohorts Participating in the CODATwins Project. <i>Twin Research and Human Genetics</i> , 2016, 19, 112-124.	0.3	21
281	Genetic predisposition to ductal carcinoma in situ of the breast. <i>Breast Cancer Research</i> , 2016, 18, 22.	2.2	43
282	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.	0.8	21
283	Multivitamin, calcium and folic acid supplements and the risk of colorectal cancer in Lynch syndrome. <i>International Journal of Epidemiology</i> , 2016, 45, 940-953.	0.9	27
284	Five endometrial cancer risk loci identified through genome-wide association analysis. <i>Nature Genetics</i> , 2016, 48, 667-674.	9.4	77
285	Anti-Allergenic hormone serum concentrations of women with germline <i>BRCA1</i> or <i>BRCA2</i> mutations. <i>Human Reproduction</i> , 2016, 31, 1126-1132.	0.4	84
286	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. <i>Cancer Research</i> , 2016, 76, 5103-5114.	0.4	100
287	Analysis of the breast cancer methylome using formalin-fixed paraffin-embedded tumour. <i>Breast Cancer Research and Treatment</i> , 2016, 160, 173-180.	1.1	6
288	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.	2.6	59

#	ARTICLE	IF	CITATIONS
289	Genome-wide measures of DNA methylation in peripheral blood and the risk of urothelial cell carcinoma: a prospective nested caseâ€“control study. <i>British Journal of Cancer</i> , 2016, 115, 664-673.	2.9	38
290	Sensitization to milk, egg and peanut from birth to 18 years: A longitudinal study of a cohort at risk of allergic disease. <i>Pediatric Allergy and Immunology</i> , 2016, 27, 83-91.	1.1	34
291	Mother's smoking and complex lung function of offspring in middle age: A cohort study from childhood. <i>Respirology</i> , 2016, 21, 911-919.	1.3	34
292	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.	1.4	33
293	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.	1.1	64
294	Use of a Novel Nonparametric Version of DEPTH to Identify Genomic Regions Associated with Prostate Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1619-1624.	1.1	7
295	rs2735383, located at a microRNA binding site in the 3â€™UTR of NBS1, is not associated with breast cancer risk. <i>Scientific Reports</i> , 2016, 6, 36874.	1.6	2
296	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, 1052-1067.	7.7	157
297	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	5.8	93
298	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breastâ€“ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	5.8	78
299	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
300	Genome-wide analysis identifies 12 loci influencing human reproductive behavior. <i>Nature Genetics</i> , 2016, 48, 1462-1472.	9.4	284
301	Reliability of DNA methylation measures from dried blood spots and mononuclear cells using the HumanMethylation450k BeadArray. <i>Scientific Reports</i> , 2016, 6, 30317.	1.6	58
302	The Charles Perkins Centre's Twins Research Node. <i>Twin Research and Human Genetics</i> , 2016, 19, 393-396.	0.3	2
303	Mammographic density assessed on paired raw and processed digital images and on paired screen-film and digital images across three mammography systems. <i>Breast Cancer Research</i> , 2016, 18, 130.	2.2	17
304	Development and External Validation of a Melanoma Risk Prediction Model Based on Self-assessed Risk Factors. <i>JAMA Dermatology</i> , 2016, 152, 889.	2.0	53
305	Age- and Tumor Subtypeâ€“Specific Breast Cancer Risk Estimates for <i>CH</i> <i>EK</i> <i>2</i> *110delC Carriers. <i>Journal of Clinical Oncology</i> , 2016, 34, 2750-2760.	0.8	152
306	Cohort Profile: The Breast Cancer Prospective Family Study Cohort (ProF-SC). <i>International Journal of Epidemiology</i> , 2016, 45, 683-692.	0.9	48

#	ARTICLE	IF	CITATIONS
307	SNP rs16906252C&gt;T Is an Expression and Methylation Quantitative Trait Locus Associated with an Increased Risk of Developing <i>MGMT</i>-Methylated Colorectal Cancer. <i>Clinical Cancer Research</i> , 2016, 22, 6266-6277.	3.2	22
308	Risk of extracolonic cancers for people with biallelic and monoallelic mutations in <i>MUTYH</i>. <i>International Journal of Cancer</i> , 2016, 139, 1557-1563.	2.3	107
309	Breast Cancer Risk Prediction Using Clinical Models and 77 Independent Risk-Associated SNPs for Women Aged Under 50 Years: Australian Breast Cancer Family Registry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 359-365.	1.1	96
310	International Consortium on Mammographic Density: Methodology and population diversity captured across 22 countries. <i>Cancer Epidemiology</i> , 2016, 40, 141-151.	0.8	19
311	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2016, 23, 77-91.	1.6	62
312	Determining the familial risk distribution of colorectal cancer: a data mining approach. <i>Familial Cancer</i> , 2016, 15, 241-251.	0.9	6
313	Variation at 2q35 (<i>PNKD</i> and <i>TMBIM1</i>) influences colorectal cancer risk and identifies a pleiotropic effect with inflammatory bowel disease. <i>Human Molecular Genetics</i> , 2016, 25, 2349-2359.	1.4	37
314	Quantifying the utility of single nucleotide polymorphisms to guide colorectal cancer screening. <i>Future Oncology</i> , 2016, 12, 503-513.	1.1	42
315	Germline mutations in <i>PMS2</i> and <i>MLH1</i> in individuals with solitary loss of PMS2 expression in colorectal carcinomas from the Colon Cancer Family Registry Cohort. <i>BMJ Open</i> , 2016, 6, e010293.	0.8	33
316	GWASeq: targeted re-sequencing follow up to GWAS. <i>BMC Genomics</i> , 2016, 17, 176.	1.2	7
317	iPreventÂ®: a tailored, web-based, decision support tool for breast cancer risk assessment and management. <i>Breast Cancer Research and Treatment</i> , 2016, 156, 171-182.	1.1	33
318	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
319	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
320	Childhood body mass index and adult mammographic density measures that predict breast cancer risk. <i>Breast Cancer Research and Treatment</i> , 2016, 156, 163-170.	1.1	19
321	Identification of Susceptibility Loci and Genes for Colorectal Cancer Risk. <i>Gastroenterology</i> , 2016, 150, 1633-1645.	0.6	97
322	Global measures of peripheral blood-derived DNA methylation as a risk factor in the development of mature B-cell neoplasms. <i>Epigenomics</i> , 2016, 8, 55-66.	1.0	35
323	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
324	Are obesity and body fat distribution associated with low back pain in women? A population-based study of 1128 Spanish twins. <i>European Spine Journal</i> , 2016, 25, 1188-1195.	1.0	50

#	ARTICLE	IF	CITATIONS
325	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	0.6	18
326	RAD51B in Familial Breast Cancer. <i>PLoS ONE</i> , 2016, 11, e0153788.	1.1	26
327	Is Chronic Low Back Pain Associated with the Prevalence of Coronary Heart Disease when Genetic Susceptibility Is Considered? A Co-Twin Control Study of Spanish Twins. <i>PLoS ONE</i> , 2016, 11, e0155194.	1.1	33
328	Methylation of Breast Cancer Predisposition Genes in Early-Onset Breast Cancer: Australian Breast Cancer Family Registry. <i>PLoS ONE</i> , 2016, 11, e0165436.	1.1	12
329	Alcohol consumption for different periods in life, intake pattern over time and all-cause mortality. <i>Journal of Public Health</i> , 2015, 37, fdu082.	1.0	20
330	Detection of skewed X-chromosome inactivation in Fragile X syndrome and X chromosome aneuploidy using quantitative melt analysis. <i>Expert Reviews in Molecular Medicine</i> , 2015, 17, e13.	1.6	12
331	Zygoty Differences in Height and Body Mass Index of Twins From Infancy to Old Age: A Study of the CODATwins Project. <i>Twin Research and Human Genetics</i> , 2015, 18, 557-570.	0.3	24
332	The CODATwins Project: The Cohort Description of Collaborative Project of Development of Anthropometrical Measures in Twins to Study Macro-Environmental Variation in Genetic and Environmental Effects on Anthropometric Traits. <i>Twin Research and Human Genetics</i> , 2015, 18, 348-360.	0.3	55
333	Genetic and Environmental Causes of Variation in the Difference Between Biological Age Based on DNA Methylation and Chronological Age for Middle-Aged Women. <i>Twin Research and Human Genetics</i> , 2015, 18, 720-726.	0.3	43
334	Quantifying the cumulative effect of low-penetrance genetic variants on breast cancer risk. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2015, 3, 182-188.	0.6	1
335	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
336	High performance computing enabling exhaustive analysis of higher order single nucleotide polymorphism interaction in Genome Wide Association Studies. <i>Health Information Science and Systems</i> , 2015, 3, S3.	3.4	24
337	Commentary: Age and frailty "not quite the same thing. <i>International Journal of Epidemiology</i> , 2015, 44, 1421-1423.	0.9	1
338	Common germline polymorphisms associated with breast cancer-specific survival. <i>Breast Cancer Research</i> , 2015, 17, 58.	2.2	26
339	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
340	Mammographic density defined by higher than conventional brightness threshold better predicts breast cancer risk for full-field digital mammograms. <i>Breast Cancer Research</i> , 2015, 17, 142.	2.2	35
341	Tools for translational epigenetic studies involving formalin-fixed paraffin-embedded human tissue: applying the Infinium HumanMethylation450 Beadchip assay to large population-based studies. <i>BMC Research Notes</i> , 2015, 8, 543.	0.6	15
342	Hormonal contraception increases risk of asthma among obese but decreases it among nonobese subjects: a prospective, population-based cohort study. <i>ERJ Open Research</i> , 2015, 1, 00026-2015.	1.1	12

#	ARTICLE	IF	CITATIONS
343	Lynch syndrome and cervical cancer. <i>International Journal of Cancer</i> , 2015, 137, 2757-2761.	2.3	13
344	SNP-SNP interaction analysis of NF- $\kappa$ B signaling pathway on breast cancer survival. <i>Oncotarget</i> , 2015, 6, 37979-37994.	0.8	20
345	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
346	Practical Problems With Clinical Guidelines for Breast Cancer Prevention Based on Remaining Lifetime Risk. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv124-djv124.	3.0	34
347	Association between Body Mass Index and Mortality for Colorectal Cancer Survivors: Overall and by Tumor Molecular Phenotype. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1229-1238.	1.1	44
348	Genetic and Environmental Variances of Bone Microarchitecture and Bone Remodeling Markers: A Twin Study. <i>Journal of Bone and Mineral Research</i> , 2015, 30, 519-527.	3.1	41
349	Association of the Colorectal CpG Island Methylator Phenotype with Molecular Features, Risk Factors, and Family History. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 512-519.	1.1	71
350	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
351	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015, 47, 373-380.	9.4	513
352	Risk of colorectal cancer for people with a mutation in both a MUTYH and a DNA mismatch repair gene. <i>Familial Cancer</i> , 2015, 14, 575-583.	0.9	11
353	Aspirin, Ibuprofen, and the Risk of Colorectal Cancer in Lynch Syndrome. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv170.	3.0	80
354	Genetic determinants of telomere length and risk of common cancers: a Mendelian randomization study. <i>Human Molecular Genetics</i> , 2015, 24, 5356-5366.	1.4	128
355	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. <i>Nature Communications</i> , 2015, 6, 7138.	5.8	138
356	A new GWAS and meta-analysis with 1000Genomes imputation identifies novel risk variants for colorectal cancer. <i>Scientific Reports</i> , 2015, 5, 10442.	1.6	109
357	Female Hormonal Factors and the Risk of Endometrial Cancer in Lynch Syndrome. <i>JAMA - Journal of the American Medical Association</i> , 2015, 314, 61.	3.8	68
358	Accuracy of Self-Reported Nevus and Pigmentation Phenotype Compared with Clinical Assessment in a Population-Based Study of Young Australian Adults. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 736-743.	1.1	15
359	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.	2.6	37
360	Identification of Novel Genetic Markers of Breast Cancer Survival. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	3.0	56

#	ARTICLE	IF	CITATIONS
361	Risk Analysis of Prostate Cancer in PRACTICAL, a Multinational Consortium, Using 25 Known Prostate Cancer Susceptibility Loci. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1121-1129.	1.1	56
362	Childhood cancers in families with and without Lynch syndrome. <i>Familial Cancer</i> , 2015, 14, 545-551.	0.9	8
363	The Heritability of Prostate Cancer—Letter. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 878-878.	1.1	8
364	Novel Associations between Common Breast Cancer Susceptibility Variants and Risk-Predicting Mammographic Density Measures. <i>Cancer Research</i> , 2015, 75, 2457-2467.	0.4	55
365	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.	9.4	357
366	A genome-wide association study for colorectal cancer identifies a risk locus in 14q23.1. <i>Human Genetics</i> , 2015, 134, 1249-1262.	1.8	28
367	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv219.	3.0	99
368	Breast cancer risk for Korean women with germline mutations in BRCA1 and BRCA2. <i>Breast Cancer Research and Treatment</i> , 2015, 152, 659-665.	1.1	18
369	Comprehensive genetic assessment of the ESR1 locus identifies a risk region for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2015, 22, 851-861.	1.6	25
370	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, 1680-1691.	1.1	24
371	SNPs and breast cancer risk prediction for African American and Hispanic women. <i>Breast Cancer Research and Treatment</i> , 2015, 154, 583-589.	1.1	49
372	Odds per Adjusted Standard Deviation: Comparing Strengths of Associations for Risk Factors Measured on Different Scales and Across Diseases and Populations: Table 1.. <i>American Journal of Epidemiology</i> , 2015, 182, 863-867.	1.6	80
373	Short-Term Risk of Colorectal Cancer in Individuals With Lynch Syndrome: A Meta-Analysis. <i>Journal of Clinical Oncology</i> , 2015, 33, 326-331.	0.8	37
374	Role of tumour molecular and pathology features to estimate colorectal cancer risk for first-degree relatives. <i>Gut</i> , 2015, 64, 101-110.	6.1	40
375	Consequences of germline variation disrupting the constitutional translational initiation codon start sites of <i>MLH1</i> and <i>BRCA2</i> : Use of potential alternative start sites and implications for predicting variant pathogenicity. <i>Molecular Carcinogenesis</i> , 2015, 54, 513-522.	1.3	14
376	Lifetime alcohol consumption and upper aero-digestive tract cancer risk in the Melbourne Collaborative Cohort Study. <i>Cancer Causes and Control</i> , 2015, 26, 297-301.	0.8	10
377	Genetic variants within the hTERT gene and the risk of colorectal cancer in Lynch syndrome. <i>Genes and Cancer</i> , 2015, 6, 445-451.	0.6	6
378	The SNP rs6500843 in 16p13.3 is associated with survival specifically among chemotherapy-treated breast cancer patients. <i>Oncotarget</i> , 2015, 6, 7390-7407.	0.8	15

#	ARTICLE	IF	CITATIONS
379	MicroRNA Related Polymorphisms and Breast Cancer Risk. <i>PLoS ONE</i> , 2014, 9, e109973.	1.1	49
380	Register4: an Australian web-enabled resource created by the National Breast Cancer Foundation to facilitate and accelerate cancer research. <i>Medical Journal of Australia</i> , 2014, 200, 460-460.	0.8	2
381	A novel colorectal cancer risk locus at 4q32.2 identified from an international genome-wide association study. <i>Carcinogenesis</i> , 2014, 35, 2512-2519.	1.3	30
382	Re: Microsatellite Instability and BRAF Mutation Testing in Colorectal Cancer Prognostication. <i>Journal of the National Cancer Institute</i> , 2014, 106, dju180-dju180.	3.0	6
383	Identification of a melanoma susceptibility locus and somatic mutation in <i>TET2</i> . <i>Carcinogenesis</i> , 2014, 35, 2097-2101.	1.3	41
384	Genetic Predisposition to In Situ and Invasive Lobular Carcinoma of the Breast. <i>PLoS Genetics</i> , 2014, 10, e1004285.	1.5	39
385	DNA Glycosylases Involved in Base Excision Repair May Be Associated with Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. <i>PLoS Genetics</i> , 2014, 10, e1004256.	1.5	47
386	2q36.3 is associated with prognosis for oestrogen receptor-negative breast cancer patients treated with chemotherapy. <i>Nature Communications</i> , 2014, 5, 4051.	5.8	16
387	Epigenome-wide methylation in DNA from peripheral blood as a marker of risk for breast cancer. <i>Breast Cancer Research and Treatment</i> , 2014, 148, 665-673.	1.1	93
388	Genetics of epilepsy. <i>Neurology</i> , 2014, 83, 1042-1048.	1.5	61
389	Prevalence and predictors of germline CDKN2A mutations for melanoma cases from Australia, Spain and the United Kingdom. <i>Hereditary Cancer in Clinical Practice</i> , 2014, 12, 20.	0.6	45
390	Cancer Risks for Relatives of Children with Cancer. <i>Journal of Cancer Epidemiology</i> , 2014, 2014, 1-4.	0.5	10
391	A Genome-wide Association Study of Early-Onset Breast Cancer Identifies <i>PFKM</i> as a Novel Breast Cancer Gene and Supports a Common Genetic Spectrum for Breast Cancer at Any Age. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 658-669.	1.1	77
392	Risk of Colorectal Cancer for Carriers of Mutations in <i>MUTYH</i> , With and Without a Family History of Cancer. <i>Gastroenterology</i> , 2014, 146, 1208-1211.e5.	0.6	180
393	Early Detection of Fragile X Syndrome: Applications of a Novel Approach for Improved Quantitative Methylation Analysis in Venous Blood and Newborn Blood Spots. <i>Clinical Chemistry</i> , 2014, 60, 963-973.	1.5	43
394	Dietary Patterns and Their Associations with Age-Related Macular Degeneration. <i>Ophthalmology</i> , 2014, 121, 1428-1434.e2.	2.5	63
395	Genome-wide association analysis identifies 11 risk variants associated with the asthma with hay fever phenotype. <i>Journal of Allergy and Clinical Immunology</i> , 2014, 133, 1564-1571.	1.5	195
396	Rare Mutations in <i>RINT1</i> Predispose Carriers to Breast and Lynch Syndrome "Spectrum Cancers. <i>Cancer Discovery</i> , 2014, 4, 804-815.	7.7	44

#	ARTICLE	IF	CITATIONS
397	Genome-wide association study identifies multiple loci associated with both mammographic density and breast cancer risk. <i>Nature Communications</i> , 2014, 5, 5303.	5.8	109
398	Associations of Mammographic Dense and Nondense Areas and Body Mass Index With Risk of Breast Cancer. <i>American Journal of Epidemiology</i> , 2014, 179, 475-483.	1.6	48
399	Should the grading of colorectal adenocarcinoma include microsatellite instability status?. <i>Human Pathology</i> , 2014, 45, 2077-2084.	1.1	44
400	Alcohol Consumption and Survival after a Breast Cancer Diagnosis: A Literature-Based Meta-analysis and Collaborative Analysis of Data for 29,239 Cases. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 934-945.	1.1	37
401	Parent-of-origin-specific allelic associations among 106 genomic loci for age at menarche. <i>Nature</i> , 2014, 514, 92-97.	13.7	548
402	Identification of susceptibility loci for colorectal cancer in a genome-wide meta-analysis. <i>Human Molecular Genetics</i> , 2014, 23, 4729-4737.	1.4	128
403	Breast-Cancer Risk in Families with Mutations in <i>PALB2</i> . <i>New England Journal of Medicine</i> , 2014, 371, 497-506.	13.9	745
404	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	5.8	105
405	Does risk of endometrial cancer for women without a germline mutation in a DNA mismatch repair gene depend on family history of endometrial cancer or colorectal cancer?. <i>Gynecologic Oncology</i> , 2014, 133, 287-292.	0.6	20
406	Cost-effectiveness of family history-based colorectal cancer screening in Australia. <i>BMC Cancer</i> , 2014, 14, 261.	1.1	24
407	Assessing and managing breast cancer risk: Clinicians' current practice and future needs. <i>Breast</i> , 2014, 23, 644-650.	0.9	44
408	Bone mineral density and the risk of breast cancer: a case-control study of Korean women. <i>Annals of Epidemiology</i> , 2014, 24, 222-227.	0.9	16
409	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.	2.2	14
410	Using SNP genotypes to improve the discrimination of a simple breast cancer risk prediction model. <i>Breast Cancer Research and Treatment</i> , 2013, 139, 887-896.	1.1	33
411	Hi-Plex for high-throughput mutation screening: application to the breast cancer susceptibility gene PALB2. <i>BMC Medical Genomics</i> , 2013, 6, 48.	0.7	13
412	Architecture of cortical bone determines in part its remodelling and structural decay. <i>Bone</i> , 2013, 55, 353-358.	1.4	31
413	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	9.4	960
414	Identification of Genetic Susceptibility Loci for Colorectal Tumors in a Genome-Wide Meta-analysis. <i>Gastroenterology</i> , 2013, 144, 799-807.e24.	0.6	292



#	ARTICLE	IF	CITATIONS
415	Australian Twin Registry: 30 Years of Progress. <i>Twin Research and Human Genetics</i> , 2013, 16, 34-42.	0.3	32
416	Fracture risk and height: An association partly accounted for by cortical porosity of relatively thinner cortices. <i>Journal of Bone and Mineral Research</i> , 2013, 28, 2017-2026.	3.1	83
417	Explaining Variance in the <i>Cumulus</i> Mammographic Measures That Predict Breast Cancer Risk: A Twins and Sisters Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 2395-2403.	1.1	36
418	Population-Based Estimate of Prostate Cancer Risk for Carriers of the HOXB13 Missense Mutation G84E. <i>PLoS ONE</i> , 2013, 8, e54727.	1.1	31
419	Inference about Causation from Examination of Familial Confounding: Application to Longitudinal Twin Data on Mammographic Density Measures that Predict Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1149-1155.	1.1	34
420	Screening Practices of Unaffected People at Familial Risk of Colorectal Cancer. <i>Cancer Prevention Research</i> , 2012, 5, 240-247.	0.7	25
421	<i>CHEK2</i>*1100delC Heterozygosity in Women With Breast Cancer Associated With Early Death, Breast Cancer-Specific Death, and Increased Risk of a Second Breast Cancer. <i>Journal of Clinical Oncology</i> , 2012, 30, 4308-4316.	0.8	162
422	Childhood Infections and the Risk of Asthma. <i>Chest</i> , 2012, 142, 647-654.	0.4	28
423	Screening practices of Australian men and women categorized as "at or slightly above average risk" of colorectal cancer. <i>Cancer Causes and Control</i> , 2012, 23, 1853-1864.	0.8	17
424	Does eczema in infancy cause hay fever, asthma, or both in childhood? Insights from a novel regression model of sibling data. <i>Journal of Allergy and Clinical Immunology</i> , 2012, 130, 1117-1122.e1.	1.5	56
425	Are genetic and environmental components of variance in mammographic density measures that predict breast cancer risk independent of within-twin pair differences in body mass index?. <i>Breast Cancer Research and Treatment</i> , 2012, 131, 553-559.	1.1	3
426	Meta-analysis of new genome-wide association studies of colorectal cancer risk. <i>Human Genetics</i> , 2012, 131, 217-234.	1.8	183
427	Remodeling markers are associated with larger intracortical surface area but smaller trabecular surface area: A twin study. <i>Bone</i> , 2011, 49, 1125-1130.	1.4	50
428	Genotype-Environment Interactions in Microsatellite Stable/Microsatellite Instability-Low Colorectal Cancer: Results from a Genome-Wide Association Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 758-766.	1.1	50
429	Metachronous colorectal cancer risk for mismatch repair gene mutation carriers "the advantage of more extensive surgery. <i>Hereditary Cancer in Clinical Practice</i> , 2011, 9, O1.	0.6	2
430	No evidence of MMTV-like env sequences in specimens from the Australian Breast Cancer Family Study. <i>Breast Cancer Research and Treatment</i> , 2011, 125, 229-235.	1.1	28
431	Disease-specific prospective family study cohorts enriched for familial risk. <i>Epidemiologic Perspectives and Innovations</i> , 2011, 8, 2.	7.0	30
432	How do women at increased, but unexplained, familial risk of breast cancer perceive and manage their risk? A qualitative interview study. <i>Hereditary Cancer in Clinical Practice</i> , 2011, 9, 7.	0.6	7

#	ARTICLE	IF	CITATIONS
433	Dependence of colorectal cancer risk on the parent-of-origin of mutations in DNA mismatch repair genes. <i>Human Mutation</i> , 2011, 32, 207-212.	1.1	9
434	The potential value of sibling controls compared with population controls for association studies of lifestyle-related risk factors: an example from the Breast Cancer Family Registry. <i>International Journal of Epidemiology</i> , 2011, 40, 1342-1354.	0.9	18
435	Bivariate Mixture Models of Serum Ferritin and Transferrin Saturation Predict Stable Components Measured 12 Years Apart in a Healthy Australian Population. <i>Blood</i> , 2011, 118, 5281-5281.	0.6	0
436	Assessing interactions between the associations of common genetic susceptibility variants, reproductive history and body mass index with breast cancer risk in the breast cancer association consortium: a combined case-control study. <i>Breast Cancer Research</i> , 2010, 12, R110.	2.2	82
437	Past recreational physical activity, body size, and all-cause mortality following breast cancer diagnosis: results from the breast cancer family registry. <i>Breast Cancer Research and Treatment</i> , 2010, 123, 531-542.	1.1	50
438	Common Breast Cancer Susceptibility Alleles and the Risk of Breast Cancer for <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers: Implications for Risk Prediction. <i>Cancer Research</i> , 2010, 70, 9742-9754.	0.4	169
439	Family History, Mammographic Density, and Risk of Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 456-463.	1.1	88
440	Influence of High-Dose Estrogen Exposure during Adolescence on Mammographic Density for Age in Adulthood. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 121-129.	1.1	7
441	Common Genetic Variants Associated with Breast Cancer and Mammographic Density Measures That Predict Disease. <i>Cancer Research</i> , 2010, 70, 1449-1458.	0.4	74
442	Is uptake of genetic testing for colorectal cancer influenced by knowledge of insurance implications?. <i>Medical Journal of Australia</i> , 2009, 191, 255-258.	0.8	58
443	<i>BRCA1</i> and <i>BRCA2</i> mutation carriers in the Breast Cancer Family Registry: an open resource for collaborative research. <i>Breast Cancer Research and Treatment</i> , 2009, 116, 379-386.	1.1	52
444	Towards more effective and equitable genetic testing for <i>BRCA1</i> and <i>BRCA2</i> mutation carriers. <i>Journal of Medical Genetics</i> , 2008, 45, 409-410.	1.5	1
445	Suggested actions from the Melbourne HVP Information Seminar. <i>Nature Precedings</i> , 2008, , .	0.1	0
446	Suggested actions from the Melbourne HVP Information Seminar. <i>Nature Precedings</i> , 2008, , .	0.1	0
447	Is childhood immunisation associated with atopic disease from age 7 to 32 years?. <i>Thorax</i> , 2007, 62, 270-275.	2.7	26
448	Colon Cancer Family Registry: An International Resource for Studies of the Genetic Epidemiology of Colon Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 2331-2343.	1.1	315
449	A common coding variant in <i>CASP8</i> is associated with breast cancer risk. <i>Nature Genetics</i> , 2007, 39, 352-358.	9.4	591
450	Genome-wide association study identifies novel breast cancer susceptibility loci. <i>Nature</i> , 2007, 447, 1087-1093.	13.7	2,165

#	ARTICLE	IF	CITATIONS
451	Rationale for, and approach to, studying modifiers of risk in persons with a genetic predisposition to colorectal cancer. <i>Current Oncology Reports</i> , 2007, 9, 202-207.	1.8	9
452	Analysis of cancer risk and BRCA1 and BRCA2 mutation prevalence in the kConFab familial breast cancer resource. <i>Breast Cancer Research</i> , 2006, 8, R12.	2.2	135
453	Tracing 8,600 participants 36 years after recruitment at age seven for the Tasmanian Asthma Study. <i>Australian and New Zealand Journal of Public Health</i> , 2006, 30, 105-110.	0.8	35
454	RESPONSE LETTER TO DR. GAU ET AL.. <i>Journal of the American Geriatrics Society</i> , 2006, 54, 1021-1022.	1.3	4
455	Determinants of Preferences for Genetic Counselling in Jewish Women. <i>Familial Cancer</i> , 2006, 5, 159-167.	0.9	8
456	Rationale for, and approach to, studying modifiers of risk in persons with a genetic predisposition to colorectal cancer. <i>Current Colorectal Cancer Reports</i> , 2006, 2, 173-178.	1.0	0
457	The Heritability of Mammographically Dense and Nondense Breast Tissue. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 612-617.	1.1	101
458	Australian Twin Registry: A Nationally Funded Resource for Medical and Scientific Research, Incorporating match and WATCH. <i>Twin Research and Human Genetics</i> , 2006, 9, 707-711.	0.3	13
459	Australian Twin Registry: a nationally funded resource for medical and scientific research, incorporating match and WATCH. <i>Twin Research and Human Genetics</i> , 2006, 9, 707-11.	0.3	3
460	Application of Genetics to the Prevention of Colorectal Cancer. , 2005, 166, 17-33.		15
461	A protein-truncating mutation in CYP17A1 in three sisters with early-onset breast cancer. <i>Human Mutation</i> , 2005, 26, 298-302.	1.1	11
462	Predictors of participation in clinical and psychosocial follow-up of the kConFab breast cancer family cohort. <i>Familial Cancer</i> , 2005, 4, 105-113.	0.9	47
463	Oral Contraceptive Use and Risk of Early-Onset Breast Cancer in Carriers and Noncarriers of BRCA1 and BRCA2 Mutations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005, 14, 350-356.	1.1	133
464	Population-based family studies in genetic epidemiology. <i>Lancet, The</i> , 2005, 366, 1397-1406.	6.3	148
465	Mammographic breast density as an intermediate phenotype for breast cancer. <i>Lancet Oncology, The</i> , 2005, 6, 798-808.	5.1	548
466	The Breast Cancer Family Registry: an infrastructure for cooperative multinational, interdisciplinary and translational studies of the genetic epidemiology of breast cancer. <i>Breast Cancer Research</i> , 2004, 6, R375-89.	2.2	255
467	Body size and composition and risk of postmenopausal breast cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2004, 13, 2117-25.	1.1	35
468	Uptake of offer to receive genetic information about BRCA1 and BRCA2 mutations in an Australian population-based study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2004, 13, 2258-63.	1.1	16

#	ARTICLE	IF	CITATIONS
469	Average age-specific cumulative risk of breast cancer according to type and site of germline mutations in BRCA1 and BRCA2 estimated from multiple-case breast cancer families attending Australian family cancer clinics. <i>Human Genetics</i> , 2003, 112, 542-551.	1.8	40
470	Regressive logistic and proportional hazards disease models for within-family analyses of measured genotypes, with application to a CYP17 polymorphism and breast cancer. <i>Genetic Epidemiology</i> , 2003, 24, 161-172.	0.6	22
471	Commentary: Case-control-family designs: a paradigm for future epidemiology research?. <i>International Journal of Epidemiology</i> , 2003, 32, 48-50.	0.9	15
472	The Australian Twin Registry. <i>Twin Research and Human Genetics</i> , 2002, 5, 329-336.	1.5	33
473	Heritability of Mammographic Density, a Risk Factor for Breast Cancer. <i>New England Journal of Medicine</i> , 2002, 347, 886-894.	13.9	537
474	More breast cancer genes?. <i>Breast Cancer Research</i> , 2001, 3, 154-7.	2.2	9
475	Association of Birth Weight and Current Body Size to Blood Pressure in Female Twins. <i>Twin Research and Human Genetics</i> , 2001, 4, 378-384.	1.5	7
476	Genetic epidemiology of female breast cancer. <i>Seminars in Cancer Biology</i> , 2001, 11, 367-374.	4.3	55
477	Association of Birth Weight and Current Body Size to Blood Pressure in Female Twins. <i>Twin Research and Human Genetics</i> , 2001, 4, 378-384.	1.5	9
478	Androgen receptor exon 1 cag repeat length and risk of ovarian cancer. <i>International Journal of Cancer</i> , 2000, 87, 637-643.	2.3	37
479	Adaptive evolution of the tumour suppressor BRCA1 in humans and chimpanzees. <i>Nature Genetics</i> , 2000, 25, 410-413.	9.4	153
480	ConFab: a research resource of Australasian breast cancer families. <i>Medical Journal of Australia</i> , 2000, 172, 463-464.	0.8	35
481	A Prospective Longitudinal Study of Serum Testosterone, Dehydroepiandrosterone Sulfate, and Sex Hormone-Binding Globulin Levels through the Menopause Transition. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2000, 85, 2832-2838.	1.8	342
482	RESPONSE: Re: HRAS1 Rare Minisatellite Alleles and Breast Cancer in Australian Women Under Age Forty Years. <i>Journal of the National Cancer Institute</i> , 2000, 92, 756-757.	3.0	2
483	Mechanisms of Bone Loss Following Allogeneic and Autologous Hemopoietic Stem Cell Transplantation. <i>Journal of Bone and Mineral Research</i> , 1999, 14, 342-350.	3.1	156
484	Likelihood-based approach to estimating twin concordance for dichotomous traits. , 1999, 16, 290-304.		76
485	Breast cancer in Australian women under the age of 40. <i>Cancer Causes and Control</i> , 1998, 9, 189-198.	0.8	101
486	Bone Mineral Density and Bone Turnover in Asthmatics Treated with Long-Term Inhaled or Oral Glucocorticoids. <i>Journal of Bone and Mineral Research</i> , 1998, 13, 1283-1289.	3.1	86

#	ARTICLE	IF	CITATIONS
487	Epilepsies in twins: Genetics of the major epilepsy syndromes. <i>Annals of Neurology</i> , 1998, 43, 435-445.	2.8	365
488	The histologic phenotypes of breast carcinoma occurring before age 40 years in women with and without BRCA1 or BRCA2 germline mutations. <i>Cancer</i> , 1998, 83, 2335-2345.	2.0	243
489	CFTR $\Delta$ F508 carrier status, risk of breast cancer before the age of 40 and histological grading in a population-based case-control study. , 1998, 79, 487-489.		19
490	The histologic phenotypes of breast carcinoma occurring before age 40 years in women with and without BRCA1 or BRCA2 germline mutations. <i>Cancer</i> , 1998, 83, 2335-2345.	2.0	4
491	GENETIC FACTORS ASSOCIATED WITH ALTERED SODIUM TRANSPORT IN HUMAN HYPERTENSION: A TWIN STUDY. <i>Clinical and Experimental Pharmacology and Physiology</i> , 1997, 24, 424-426.	0.9	2
492	Changes in physical activity and health outcomes in a population-based cohort of mid-life Australian-born women. <i>Australian and New Zealand Journal of Public Health</i> , 1997, 21, 682-687.	0.8	20
493	AT-tributable risks?. <i>Nature Genetics</i> , 1997, 15, 226-226.	9.4	56
494	Regressive logistic modeling of familial aggregation for asthma in 7,394 population-based nuclear families. <i>Genetic Epidemiology</i> , 1997, 14, 317-332.	0.6	61
495	The outgoing, the rebellious and the anxious: Are adolescent personality dimensions related to the uptake of smoking?. <i>Psychology and Health</i> , 1996, 12, 73-85.	1.2	9
496	Familial temporal lobe epilepsy: A common disorder identified in twins. <i>Annals of Neurology</i> , 1996, 40, 227-235.	2.8	211
497	Some public health issues in the current state of genetic testing for breast cancer in Australia. <i>Australian and New Zealand Journal of Public Health</i> , 1996, 20, 467-472.	0.8	9
498	Retired elite female ballet dancers and nonathletic controls have similar bone mineral density at weightbearing sites. <i>Journal of Bone and Mineral Research</i> , 1996, 11, 1566-1574.	3.1	53
499	Risk factors and preventive strategies for breast cancer. <i>Medical Journal of Australia</i> , 1995, 163, 435-440.	0.8	12
500	Do selected kinanthropometric and performance variables predict injuries in female netball players?. <i>Journal of Sports Sciences</i> , 1995, 13, 213-222.	1.0	57
501	Determinants of bone mass in 10- to 26-year-old females: A twin study. <i>Journal of Bone and Mineral Research</i> , 1995, 10, 558-567.	3.1	160
502	Bone density determinants in elderly women: A twin study. <i>Journal of Bone and Mineral Research</i> , 1995, 10, 1607-1613.	3.1	181
503	Increase in the self-reported prevalence of asthma and hay fever in adults over the last generation: a matched parent-offspring study. <i>Australian Journal of Public Health</i> , 1995, 19, 120-124.	0.2	56
504	Genetic factors in alcohol use-a genetic epidemiological perspective. <i>Drug and Alcohol Review</i> , 1994, 13, 375-384.	1.1	0

#	ARTICLE	IF	CITATIONS
505	A MULTIVARIATE NORMAL MODEL FOR PEDIGREE AND LONGITUDINAL DATA AND THE SOFTWARE "FISHER". The Australian Journal of Statistics, 1994, 36, 153-176.	0.2	46
506	Reduced femoral neck bone density in the daughters of women with hip fractures: The role of low peak bone density in the pathogenesis of osteoporosis. Journal of Bone and Mineral Research, 1994, 9, 739-743.	3.1	115
507	Factors in childhood as predictors of asthma in adult life. BMJ: British Medical Journal, 1994, 309, 90-93.	2.4	224
508	The associations between childhood asthma and atopy, and parental asthma, hay fever and smoking. Paediatric and Perinatal Epidemiology, 1993, 7, 67-76.	0.8	58
509	Changes in axial bone density with age: A twin study. Journal of Bone and Mineral Research, 1993, 8, 11-17.	3.1	168
510	Familial Aggregation of a Disease Consequent upon Correlation between Relatives in a Risk Factor Measured on a Continuous Scale. American Journal of Epidemiology, 1992, 136, 1138-1147.	1.6	118
511	Pedigree analysis of blood pressure in subjects from rural Greece and relatives who migrated to Melbourne, Australia. Genetic Epidemiology, 1992, 9, 225-238.	0.6	12
512	Costs and benefits of the use of commercial market research approaches in large scale surveys. Medical Journal of Australia, 1992, 157, 504-504.	0.8	14
513	PROGNOSIS OF MALE PATIENTS WITH TREATED HYPERTENSION FOLLOWED OVER 15 YEARS. Clinical and Experimental Pharmacology and Physiology, 1990, 17, 211-213.	0.9	2
514	A family study of panic disorder: Reanalysis using a regressive logistic model that incorporates a sibship environment. Genetic Epidemiology, 1990, 7, 151-161.	0.6	13
515	Twin concordance for a binary trait: III. A bivariate analysis of hay fever and asthma. Genetic Epidemiology, 1990, 7, 277-289.	0.6	44
516	Modelling sibship environment in the regressive logistic model for familial disease. Genetic Epidemiology, 1989, 6, 235-240.	0.6	8
517	A random walk model for evaluating clinical trials involving serial observations. Statistics in Medicine, 1988, 7, 581-590.	0.8	8
518	Enalapril & Nifedipine in Essential Hypertension: Synergism of the Hypotensive Effects in Combination. Clinical and Experimental Hypertension, 1988, 10, 779-789.	0.3	11
519	A family study of panic disorder. Genetic Epidemiology, 1987, 4, 33-41.	0.6	36
520	A log-linear model for binary pedigree data. Genetic Epidemiology, 1986, 3, 73-82.	0.6	25
521	THE CONTRIBUTION OF W. H. ARCHER TO VITAL STATISTICS IN THE COLONY OF VICTORIA. The Australian Journal of Statistics, 1986, 28, 124-137.	0.2	10
522	Factors that Determine the Response of People with Mild Hypertension to A Reduced Sodium Intake. Clinical and Experimental Hypertension, 1986, 8, 941-962.	0.3	11

#	ARTICLE	IF	CITATIONS
523	A genetic and environmental analysis of a twin family study of alcohol use, anxiety, and depression. <i>Genetic Epidemiology</i> , 1984, 1, 63-79.	0.6	99
524	The Utility of a Multivariate Normal Model for Studying Familial Patterns in Medical and Psychiatric Data. <i>Australian and New Zealand Journal of Psychiatry</i> , 1983, 17, 342-348.	1.3	4
525	EXTENSIONS TO MULTIVARIATE NORMAL MODELS FOR PEDIGREE ANALYSIS: II. MODELING THE EFFECT OF SHARED ENVIRONMENT IN THE ANALYSIS OF VARIATION IN BLOOD LEAD LEVELS. <i>American Journal of Epidemiology</i> , 1983, 117, 344-355.	1.6	56
526	Twin Concordance for a Binary Trait. <i>Statistical Models Illustrated With Data on Drinking Status</i> . <i>Acta Geneticae Medicae Et Gemellologiae</i> , 1983, 32, 127-137.	0.2	41
527	GENETIC ANALYSIS OF SYSTOLIC BLOOD PRESSURE IN MELBOURNE FAMILIES. <i>Clinical and Experimental Pharmacology and Physiology</i> , 1982, 9, 247-252.	0.9	18
528	Analysis of variation in blood lead levels in Melbourne families. <i>Medical Journal of Australia</i> , 1982, 2, 573-576.	0.8	15
529	Suggested actions from the Melbourne HVP Information Seminar. <i>Nature Precedings</i> , 0, , .	0.1	1
530	Improved definition of growing pains: A common familial primary pain disorder of early childhood. <i>Paediatric and Neonatal Pain</i> , 0, , .	0.6	1
531	Women's thoughts on receiving and sharing genetic information: Considerations for genetic counseling. <i>Journal of Genetic Counseling</i> , 0, , .	0.9	1