

# Janet E Olson

## List of Publications by Year in Descending Order

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**Version:** 2024-04-27

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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.

130  
papers

9,693  
citations

40  
h-index

98  
g-index

143  
ext. papers

11,802  
ext. citations

8.8  
avg, IF

4.38  
L-index

#	Paper	IF	Citations
130	Rare germline copy number variants (CNVs) and breast cancer risk.. <i>Communications Biology</i> , <b>2022</b> , 5, 65	6.7	0
129	A hybrid model to identify fall occurrence from electronic health records.. <i>International Journal of Medical Informatics</i> , <b>2022</b> , 162, 104736	5.3	1
128	Genome-wide and transcriptome-wide association studies of mammographic density phenotypes reveal novel loci.. <i>Breast Cancer Research</i> , <b>2022</b> , 24, 27	8.3	1
127	Genome-wide interaction analysis of menopausal hormone therapy use and breast cancer risk among 62,370 women.. <i>Scientific Reports</i> , <b>2022</b> , 12, 6199	4.9	
126	Prevalence and Overall Survival of Low Count Monoclonal B-Cell Lymphocytosis (LC-MBL): A Screening Study of 8,297 Individuals from the Mayo Clinic Biobank. <i>Blood</i> , <b>2021</b> , 138, 2632-2632	2.2	0
125	Relationship and Susceptibility to Serious Infections Among Monoclonal B-Cell Lymphocytosis (MBL), Monoclonal Gammopathy of Undetermined Significance (MGUS), and Clonal Hematopoiesis (CH) Premalignant Conditions. <i>Blood</i> , <b>2021</b> , 138, 3739-3739	2.2	
124	Genetic Predictors of Chemotherapy-Induced Peripheral Neuropathy from Paclitaxel, Carboplatin and Oxaliplatin: NCCTG/Alliance N08C1, N08CA and N08CB Study. <i>Cancers</i> , <b>2021</b> , 13,	6.6	6
123	Genetic basis of hypercholesterolemia in adults. <i>Npj Genomic Medicine</i> , <b>2021</b> , 6, 28	6.2	4
122	Pathway to Ascertain the Role of Pharmacogenomics in Healthcare Utilization Outcomes [Response to Letter]. <i>Pharmacogenomics and Personalized Medicine</i> , <b>2021</b> , 14, 545-546	2.1	
121	Longitudinal cohorts for harnessing the electronic health record for disease prediction in a US population. <i>BMJ Open</i> , <b>2021</b> , 11, e044353	3	2
120	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> , <b>2021</b> , 108, 1190-1203	11	1
119	Risk of serious infection among individuals with and without low count monoclonal B-cell lymphocytosis (MBL). <i>Leukemia</i> , <b>2021</b> , 35, 239-244	10.7	7
118	Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , <b>2021</b> , 113, 329-337	9.7	14
117	CYP3A7*1C allele: linking premenopausal oestrone and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , <b>2021</b> , 124, 842-854	8.7	2
116	Real-World Experiences With Yoga on Cancer-Related Symptoms in Women With Breast Cancer. <i>Global Advances in Health and Medicine</i> , <b>2021</b> , 10, 2164956120984140	1.9	4
115	Genetic Variations and Health-Related Quality of Life (HRQOL): A Genome-Wide Study Approach. <i>Cancers</i> , <b>2021</b> , 13,	6.6	1
114	A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. <i>Nature Communications</i> , <b>2021</b> , 12, 1078	17.4	4

113	A Population-Based Study of Genes Previously Implicated in Breast Cancer. <i>New England Journal of Medicine</i> , <b>2021</b> , 384, 440-451	59.2	115
112	Impact of Pharmacogenomic Information on Values of Care and Quality of Life Associated with Codeine and Tramadol-Related Adverse Drug Events. <i>Mayo Clinic Proceedings Innovations, Quality &amp; Outcomes</i> , <b>2021</b> , 5, 35-45	3.1	1
111	No Association Between Pharmacogenomics Variants and Hospital and Emergency Department Utilization: A Mayo Clinic Biobank Retrospective Study. <i>Pharmacogenomics and Personalized Medicine</i> , <b>2021</b> , 14, 229-237	2.1	2
110	Polygenic risk score and risk of monoclonal B-cell lymphocytosis in caucasians and risk of chronic lymphocytic leukemia (CLL) in African Americans. <i>Leukemia</i> , <b>2021</b> ,	10.7	1
109	Risk of Breast Cancer Among Carriers of Pathogenic Variants in Breast Cancer Predisposition Genes Varies by Polygenic Risk Score. <i>Journal of Clinical Oncology</i> , <b>2021</b> , 39, 2564-2573	2.2	12
108	Ascertainment of delirium status using natural language processing from electronic health records. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , <b>2020</b> ,	6.4	3
107	Assessing the stability of biobank donor preferences regarding sample use: evidence supporting the value of dynamic consent. <i>European Journal of Human Genetics</i> , <b>2020</b> , 28, 1168-1177	5.3	5
106	Inferring multimodal latent topics from electronic health records. <i>Nature Communications</i> , <b>2020</b> , 11, 2536	17.4	13
105	Colorectal cancer outcomes after screening with the multi-target stool DNA assay: protocol for a large-scale, prospective cohort study (the Voyage study). <i>BMJ Open Gastroenterology</i> , <b>2020</b> , 7, e000353	3.9	3
104	An assessment of patient perspectives on pharmacogenomics educational materials. <i>Pharmacogenomics</i> , <b>2020</b> , 21, 347-358	2.6	3
103	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , <b>2020</b> , 10, 9688	4.9	2
102	Impact of Diverse Data Sources on Computational Phenotyping. <i>Frontiers in Genetics</i> , <b>2020</b> , 11, 556	4.5	1
101	Evaluation of Germline Genetic Testing Criteria in a Hospital-Based Series of Women With Breast Cancer. <i>Journal of Clinical Oncology</i> , <b>2020</b> , 38, 1409-1418	2.2	28
100	Characteristics Associated With Recruitment and Re-contact in Mayo Clinic Biobank. <i>Frontiers in Public Health</i> , <b>2020</b> , 8, 9	6	2
99	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , <b>2020</b> , 11, 312	17.4	20
98	Real-world experiences with acupuncture among breast cancer survivors: a cross-sectional survey study. <i>Supportive Care in Cancer</i> , <b>2020</b> , 28, 5833-5838	3.9	3
97	Polygenic Risk Score and Risk of Chronic Lymphocytic Leukemia, Monoclonal B-Cell Lymphocytosis (MBL), and MBL Subtypes. <i>Blood</i> , <b>2020</b> , 136, 35-36	2.2	
96	N-terminal pro-brain natriuretic peptide levels after receipt of anthracycline for breast cancer.. <i>Journal of Clinical Oncology</i> , <b>2020</b> , 38, e24103-e24103	2.2	

95	Challenges in returning results in a genomic medicine implementation study: the Return of Actionable Variants Empirical (RAVE) study. <i>Npj Genomic Medicine</i> , <b>2020</b> , 5, 19	6.2	2
94	Mini-Review of Laboratory Operations in Biobanking: Building Biobanking Resources for Translational Research. <i>Frontiers in Public Health</i> , <b>2020</b> , 8, 362	6	1
93	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , <b>2020</b> , 107, 837-848	11	12
92	Association of Genetic Variants at With Chemotherapy-Related Heart Failure. <i>Frontiers in Cardiovascular Medicine</i> , <b>2020</b> , 7, 142	5.4	3
91	Cohort Profile: The Right Drug, Right Dose, Right Time: Using Genomic Data to Individualize Treatment Protocol (RIGHT Protocol). <i>International Journal of Epidemiology</i> , <b>2020</b> , 49, 23-24k	7.8	19
90	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , <b>2019</b> , 9, 12524	4.9	2
89	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> , <b>2019</b> , 21, 68	8.3	18
88	Managing the Unimaginable: Biobank Participant Views on Reconsent for Whole Genome Sequencing of Stored Biospecimens. <i>Biopreservation and Biobanking</i> , <b>2019</b> , 17, 296-302	2.1	4
87	Long-term neurotoxicity in women with breast cancer.. <i>Journal of Clinical Oncology</i> , <b>2019</b> , 37, e23089-e23089	0	0
86	Acupuncture: Real-world patient-reported outcomes of treatment-related symptoms in breast cancer survivors.. <i>Journal of Clinical Oncology</i> , <b>2019</b> , 37, e23111-e23111	2.2	0
85	Characteristics and utilisation of the Mayo Clinic Biobank, a clinic-based prospective collection in the USA: cohort profile. <i>BMJ Open</i> , <b>2019</b> , 9, e032707	3	9
84	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , <b>2019</b> , 104, 21-34	11	363
83	The BRCA2 c.68-7T>A variant is not pathogenic: A model for clinical calibration of spliceogenicity. <i>Human Mutation</i> , <b>2018</b> , 39, 729-741	4.7	16
82	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> , <b>2018</b> , 8, 6574	4.9	19
81	Germline Rare Coding Variants and Risk of Pancreatic Cancer in Minority Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2018</b> , 27, 1364-1370	4	13
80	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , <b>2018</b> , 50, 968-978	36.3	101
79	Association between the Risk of Low/High-Count Monoclonal B-Cell Lymphocytosis (MBL) and the Chronic Lymphocytic Leukemia (CLL) Polygenic Risk Score (PRS). <i>Blood</i> , <b>2018</b> , 132, 5538-5538	2.2	0
78	Association between an individual housing-based socioeconomic index and inconsistent self-reporting of health conditions: a prospective cohort study in the Mayo Clinic Biobank. <i>BMJ Open</i> , <b>2018</b> , 8, e020054	3	6

77	The Return of Actionable Variants Empirical (RAVE) Study, a Mayo Clinic Genomic Medicine Implementation Study: Design and Initial Results. <i>Mayo Clinic Proceedings</i> , <b>2018</b> , 93, 1600-1610	6.4	20
76	Developing a Process for Returning Medically Actionable Genomic Variants to Latino Patients in a Federally Qualified Health Center. <i>Public Health Genomics</i> , <b>2018</b> , 21, 77-84	1.9	14
75	Association of mitochondrial DNA copy number with self-rated health status. <i>The Application of Clinical Genetics</i> , <b>2018</b> , 11, 121-127	3.1	5
74	Ethical Considerations Related to Return of Results from Genomic Medicine Projects: The eMERGE Network (Phase III) Experience. <i>Journal of Personalized Medicine</i> , <b>2018</b> , 8,	3.6	32
73	Participant-perceived understanding and perspectives on pharmacogenomics: the Mayo Clinic RIGHT protocol (Right Drug, Right Dose, Right Time). <i>Genetics in Medicine</i> , <b>2017</b> , 19, 819-825	8.1	35
72	Variability in assigning pathogenicity to incidental findings: insights from LDLR sequence linked to the electronic health record in 1013 individuals. <i>European Journal of Human Genetics</i> , <b>2017</b> , 25, 410-415	5.3	7
71	Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , <b>2017</b> , 77, 2789-2799	10.1	49
70	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , <b>2017</b> , 551, 92-94	50.4	643
69	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , <b>2017</b> , 49, 1767-1778	36.3	186
68	Association between Alcohol Consumption, Folate Intake, and Risk of Pancreatic Cancer: A Case-Control Study. <i>Nutrients</i> , <b>2017</b> , 9,	6.7	6
67	Individual housing-based socioeconomic status predicts risk of accidental falls among adults. <i>Annals of Epidemiology</i> , <b>2017</b> , 27, 415-420.e2	6.4	24
66	Multidisciplinary model to implement pharmacogenomics at the point of care. <i>Genetics in Medicine</i> , <b>2017</b> , 19, 421-429	8.1	54
65	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , <b>2017</b> , 19, 599-603	6.3	51
64	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , <b>2017</b> , 46, 1814-1822	7.8	27
63	Zoledronic acid for treatment of osteopenia and osteoporosis in women with primary breast cancer undergoing adjuvant aromatase inhibitor therapy: a 5-year follow-up. <i>Supportive Care in Cancer</i> , <b>2016</b> , 24, 1219-26	3.9	12
62	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , <b>2016</b> , 141, 386-401	4.9	15
61	An intergenic risk locus containing an enhancer deletion in 2q35 modulates breast cancer risk by deregulating IGFBP5 expression. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 3863-3876	5.6	24
60	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , <b>2016</b> , 7, 11375	17.4	64

59	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , <b>2016</b> , 7, 12675	17.4	53
58	Personalizing Aspirin Use for Targeted Breast Cancer Chemoprevention in Postmenopausal Women. <i>Mayo Clinic Proceedings</i> , <b>2016</b> , 91, 71-80	6.4	15
57	Combined genetic and splicing analysis of BRCA1 c.[594-2A>C; 641A>G] highlights the relevance of naturally occurring in-frame transcripts for developing disease gene variant classification algorithms. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 2256-2268	5.6	55
56	A novel housing-based socioeconomic measure predicts hospitalisation and multiple chronic conditions in a community population. <i>Journal of Epidemiology and Community Health</i> , <b>2016</b> , 70, 286-91	5.1	29
55	Improvement in Cardiovascular Risk Prediction with Electronic Health Records. <i>Journal of Cardiovascular Translational Research</i> , <b>2016</b> , 9, 214-222	3.3	27
54	Incorporating a Genetic Risk Score Into Coronary Heart Disease Risk Estimates: Effect on Low-Density Lipoprotein Cholesterol Levels (the MI-GENES Clinical Trial). <i>Circulation</i> , <b>2016</b> , 133, 1181-8	16.7	138
53	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , <b>2016</b> , 48, 374-86	36.3	93
52	Heterogeneity of luminal breast cancer characterised by immunohistochemical expression of basal markers. <i>British Journal of Cancer</i> , <b>2016</b> , 114, 298-304	8.7	5
51	RAD51B in Familial Breast Cancer. <i>PLoS ONE</i> , <b>2016</b> , 11, e0153788	3.7	18
50	Impact of demographics on human gut microbial diversity in a US Midwest population. <i>PeerJ</i> , <b>2016</b> , 4, e1514	3.1	51
49	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> , <b>2016</b> , 13, e1002105	11.6	80
48	Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. <i>International Journal of Cancer</i> , <b>2016</b> , 139, 1303-1317	7.5	26
47	Aspirin use and the risk of cholangiocarcinoma. <i>Hepatology</i> , <b>2016</b> , 64, 785-96	11.2	63
46	PALB2, CHEK2 and ATM rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , <b>2016</b> , 53, 800-811	5.8	121
45	Genes associated with histopathologic features of triple negative breast tumors predict molecular subtypes. <i>Breast Cancer Research and Treatment</i> , <b>2016</b> , 157, 117-31	4.4	17
44	Genetic predisposition to ductal carcinoma in situ of the breast. <i>Breast Cancer Research</i> , <b>2016</b> , 18, 22	8.3	31
43	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. <i>Cancer Causes and Control</i> , <b>2016</b> , 27, 679-93	2.8	15
42	Integrating Pharmacogenomics into Clinical Practice: Promise vs Reality. <i>American Journal of Medicine</i> , <b>2016</b> , 129, 1093-1099.e1	2.4	42

41	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> , <b>2015</b> , 36, 256-71	4.6	12
40	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , <b>2015</b> , 47, 373-80	36.3	406
39	The contributions of breast density and common genetic variation to breast cancer risk. <i>Journal of the National Cancer Institute</i> , <b>2015</b> , 107,	9.7	128
38	A Robust e-Epidemiology Tool in Phenotyping Heart Failure with Differentiation for Preserved and Reduced Ejection Fraction: the Electronic Medical Records and Genomics (eMERGE) Network. <i>Journal of Cardiovascular Translational Research</i> , <b>2015</b> , 8, 475-83	3.3	25
37	Identification of novel genetic markers of breast cancer survival. <i>Journal of the National Cancer Institute</i> , <b>2015</b> , 107,	9.7	38
36	Novel Associations between Common Breast Cancer Susceptibility Variants and Risk-Predicting Mammographic Density Measures. <i>Cancer Research</i> , <b>2015</b> , 75, 2457-67	10.1	45
35	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. <i>Nature Genetics</i> , <b>2015</b> , 47, 1294-1303	36.3	226
34	Annexin A1 expression in a pooled breast cancer series: association with tumor subtypes and prognosis. <i>BMC Medicine</i> , <b>2015</b> , 13, 156	11.4	37
33	Design of a randomized controlled trial of disclosing genomic risk of coronary heart disease: the Myocardial Infarction Genes (MI-GENES) study. <i>BMC Medical Genomics</i> , <b>2015</b> , 8, 51	3.7	13
32	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. <i>Journal of the National Cancer Institute</i> , <b>2015</b> , 107,	9.7	74
31	Fine-scale mapping of the 4q24 locus identifies two independent loci associated with breast cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 1680-91	4	17
30	A comprehensive evaluation of interaction between genetic variants and use of menopausal hormone therapy on mammographic density. <i>Breast Cancer Research</i> , <b>2015</b> , 17, 110	8.3	13
29	How well do whole exome sequencing results correlate with medical findings? A study of 89 Mayo Clinic Biobank samples. <i>Frontiers in Genetics</i> , <b>2015</b> , 6, 244	4.5	8
28	SNP-SNP interaction analysis of NF- $\kappa$ B signaling pathway on breast cancer survival. <i>Oncotarget</i> , <b>2015</b> , 6, 37979-94	3.3	19
27	Prediction of breast cancer risk based on profiling with common genetic variants. <i>Journal of the National Cancer Institute</i> , <b>2015</b> , 107,	9.7	324
26	Inherited mutations in 17 breast cancer susceptibility genes among a large triple-negative breast cancer cohort unselected for family history of breast cancer. <i>Journal of Clinical Oncology</i> , <b>2015</b> , 33, 304-11 <sup>2</sup>	4.6	435
25	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , <b>2015</b> , 24, 2966-84	5.6	36
24	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> , <b>2015</b> , 96, 5-20	11	59

23	The SNP rs6500843 in 16p13.3 is associated with survival specifically among chemotherapy-treated breast cancer patients. <i>Oncotarget</i> , <b>2015</b> , 6, 7390-407	3.3	14
22	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , <b>2014</b> , 4, 4999	17.4	87
21	Genetic variation at CYP3A is associated with age at menarche and breast cancer risk: a case-control study. <i>Breast Cancer Research</i> , <b>2014</b> , 16, R51	8.3	12
20	MicroRNA related polymorphisms and breast cancer risk. <i>PLoS ONE</i> , <b>2014</b> , 9, e109973	3.7	37
19	Validation of a population coronary disease predictive system: the CASSANDRA model. <i>Journal of Epidemiology and Community Health</i> , <b>2014</b> , 68, 1009	5.1	
18	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 6096-111	5.6	48
17	Refined histopathological predictors of BRCA1 and BRCA2 mutation status: a large-scale analysis of breast cancer characteristics from the BCAC, CIMBA, and ENIGMA consortia. <i>Breast Cancer Research</i> , <b>2014</b> , 16, 3419	8.3	82
16	Preemptive genotyping for personalized medicine: design of the right drug, right dose, right time-using genomic data to individualize treatment protocol. <i>Mayo Clinic Proceedings</i> , <b>2014</b> , 89, 25-33	6.4	213
15	Performance of automated scoring of ER, PR, HER2, CK5/6 and EGFR in breast cancer tissue microarrays in the Breast Cancer Association Consortium <b>2014</b> , n/a-n/a		1
14	Fine-scale mapping of the FGFR2 breast cancer risk locus: putative functional variants differentially bind FOXA1 and E2F1. <i>American Journal of Human Genetics</i> , <b>2013</b> , 93, 1046-60	11	80
13	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , <b>2013</b> , 45, 371-84, 384e1-2	36.3	422
12	The Mayo Clinic Biobank: a building block for individualized medicine. <i>Mayo Clinic Proceedings</i> , <b>2013</b> , 88, 952-62	6.4	144
11	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , <b>2013</b> , 45, 392-8, 398e1-2	36.3	327
10	CHEK2*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> , <b>2012</b> , 30, 4308-16	2.2	134
9	Heterogeneity of breast cancer associations with five susceptibility loci by clinical and pathological characteristics. <i>PLoS Genetics</i> , <b>2008</b> , 4, e1000054	6	280
8	A common coding variant in CASP8 is associated with breast cancer risk. <i>Nature Genetics</i> , <b>2007</b> , 39, 352-8, 356e1-2	36.3	557
7	Genome-wide association study identifies novel breast cancer susceptibility loci. <i>Nature</i> , <b>2007</b> , 447, 1087-93	59.4	1957
6	Postmenopausal cancer risk after self-reported endometriosis diagnosis in the Iowa Women's Health Study. <i>Cancer</i> , <b>2002</b> , 94, 1612-8	6.4	105



5	The continuing increase in the incidence of primary central nervous system non-Hodgkin lymphoma: a surveillance, epidemiology, and end results analysis. <i>Cancer</i> , <b>2002</b> , 95, 1504-10	6.4	251
4	Perception of genetic risk among genetic counselors. <i>Journal of Genetic Counseling</i> , <b>2000</b> , 9, 47-59	2.5	7
3	An investigation of the biological basis of an interaction of abdominal fat distribution and family history of breast cancer. A nested study of sisters in the Iowa Women's Health Study (United States). <i>Cancer Causes and Control</i> , <b>2000</b> , 11, 941-54	2.8	3
2	Does a family history of cancer increase the risk for postmenopausal endometrial carcinoma? <b>1999</b> , 85, 2444-2449		16
1	Does a family history of cancer increase the risk for postmenopausal endometrial carcinoma? <b>1999</b> , 85, 2444		2