

Jan A Lubiński

List of Publications by Year in descending order

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Version: 2024-02-01

535
papers

42,316
citations

2423

97
h-index

3714

179
g-index

556
all docs

556
docs citations

556
times ranked

39874
citing authors

#	ARTICLE	IF	CITATIONS
1	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099
2	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. <i>American Journal of Human Genetics</i> , 2015, 97, 576-592.	2.6	1,098
3	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	9.4	960
4	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. <i>Nature</i> , 2022, 604, 502-508.	13.7	929
5	Poly(ADP-Ribose) Polymerase Inhibition: Frequent Durable Responses in <i>BRCA</i> Carrier Ovarian Cancer Correlating With Platinum-Free Interval. <i>Journal of Clinical Oncology</i> , 2010, 28, 2512-2519.	0.8	877
6	Long-term effect of aspirin on cancer risk in carriers of hereditary colorectal cancer: an analysis from the CAPP2 randomised controlled trial. <i>Lancet</i> , The, 2011, 378, 2081-2087.	6.3	849
7	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
8	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. <i>Nature Genetics</i> , 2018, 50, 928-936.	9.4	652
9	Associations of Breast Cancer Risk Factors With Tumor Subtypes: A Pooled Analysis From the Breast Cancer Association Consortium Studies. <i>Journal of the National Cancer Institute</i> , 2011, 103, 250-263.	3.0	596
10	Partitioning Heritability of Regulatory and Cell-Type-Specific Variants across 11 Common Diseases. <i>American Journal of Human Genetics</i> , 2014, 95, 535-552.	2.6	569
11	Salpingo-oophorectomy and the Risk of Ovarian, Fallopian Tube, and Peritoneal Cancers in Women With a <i>BRCA1</i> or <i>BRCA2</i> Mutation. <i>JAMA - Journal of the American Medical Association</i> , 2006, 296, 185.	3.8	544
12	Breast Cancer Risk Genes Association Analysis in More than 113,000 Women. <i>New England Journal of Medicine</i> , 2021, 384, 428-439.	13.9	532
13	Impact of Oophorectomy on Cancer Incidence and Mortality in Women With a <i>BRCA1</i> or <i>BRCA2</i> Mutation. <i>Journal of Clinical Oncology</i> , 2014, 32, 1547-1553.	0.8	523
14	Lung cancer susceptibility locus at 5p15.33. <i>Nature Genetics</i> , 2008, 40, 1404-1406.	9.4	514
15	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
16	Pathologic Complete Response Rates in Young Women With <i>BRCA1</i> -Positive Breast Cancers After Neoadjuvant Chemotherapy. <i>Journal of Clinical Oncology</i> , 2010, 28, 375-379.	0.8	500
17	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384.	9.4	493
18	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. <i>Nature Genetics</i> , 2013, 45, 385-391.	9.4	492

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19	Phase II, Open-Label, Randomized, Multicenter Study Comparing the Efficacy and Safety of Olaparib, a Poly (ADP-Ribose) Polymerase Inhibitor, and Pegylated Liposomal Doxorubicin in Patients With <i>BRCA1</i> or <i>BRCA2</i> Mutations and Recurrent Ovarian Cancer. <i>Journal of Clinical Oncology</i> , 2012, 30, 372-379.	0.8	445
20	Integrated Proteogenomic Characterization of Clear Cell Renal Cell Carcinoma. <i>Cell</i> , 2019, 179, 964-983.e31.	13.5	430
21	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
22	Genome-wide association study identifies three loci associated with melanoma risk. <i>Nature Genetics</i> , 2009, 41, 920-925.	9.4	422
23	Breast Cancer Risk Following Bilateral Oophorectomy in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers: An International Case-Control Study. <i>Journal of Clinical Oncology</i> , 2005, 23, 7491-7496.	0.8	408
24	Association of Type and Location of <i>BRCA1</i> and <i>BRCA2</i> Mutations With Risk of Breast and Ovarian Cancer. <i>JAMA - Journal of the American Medical Association</i> , 2015, 313, 1347.	3.8	390
25	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	9.4	374
26	Rare variants of large effect in <i>BRCA2</i> and <i>CHEK2</i> affect risk of lung cancer. <i>Nature Genetics</i> , 2014, 46, 736-741.	9.4	360
27	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and <i>BRCA1</i> -mediated DNA repair. <i>Nature Genetics</i> , 2015, 47, 1294-1303.	9.4	357
28	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 362-370.	9.4	326
29	A genome-wide association study identifies susceptibility loci for ovarian cancer at 2q31 and 8q24. <i>Nature Genetics</i> , 2010, 42, 874-879.	9.4	321
30	Oral Contraceptives and the Risk of Breast Cancer in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>Journal of the National Cancer Institute</i> , 2002, 94, 1773-1779.	3.0	318
31	A locus on 19p13 modifies risk of breast cancer in <i>BRCA1</i> mutation carriers and is associated with hormone receptor-negative breast cancer in the general population. <i>Nature Genetics</i> , 2010, 42, 885-892.	9.4	309
32	International variation in rates of uptake of preventive options in <i>BRCA1</i> and <i>BRCA2</i> mutation carriers. <i>International Journal of Cancer</i> , 2008, 122, 2017-2022.	2.3	306
33	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
34	Genetic Structure of Europeans: A View from the North-East. <i>PLoS ONE</i> , 2009, 4, e5472.	1.1	279
35	A genome-wide association study identifies a new ovarian cancer susceptibility locus on 9p22.2. <i>Nature Genetics</i> , 2009, 41, 996-1000.	9.4	276
36	Effect of Aspirin or Resistant Starch on Colorectal Neoplasia in the Lynch Syndrome. <i>New England Journal of Medicine</i> , 2008, 359, 2567-2578.	13.9	273

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37	Contribution of Germline Mutations in the <i>RAD51B</i> , <i>RAD51C</i> , and <i>RAD51D</i> Genes to Ovarian Cancer in the Population. <i>Journal of Clinical Oncology</i> , 2015, 33, 2901-2907.	0.8	266
38	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. <i>Nature Genetics</i> , 2011, 43, 785-791.	9.4	265
39	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
40	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. <i>Nature Genetics</i> , 2021, 53, 65-75.	9.4	264
41	Dose-Response Association of CD8 ⁺ Tumor-Infiltrating Lymphocytes and Survival Time in High-Grade Serous Ovarian Cancer. <i>JAMA Oncology</i> , 2017, 3, e173290.	3.4	260
42	Genome-wide association analysis identifies three new breast cancer susceptibility loci. <i>Nature Genetics</i> , 2012, 44, 312-318.	9.4	256
43	Tamoxifen and contralateral breast cancer in <i>BRCA1</i> and <i>BRCA2</i> carriers: An update. <i>International Journal of Cancer</i> , 2006, 118, 2281-2284.	2.3	246
44	Genome-Wide Association Study in <i>BRCA1</i> Mutation Carriers Identifies Novel Loci Associated with Breast and Ovarian Cancer Risk. <i>PLoS Genetics</i> , 2013, 9, e1003212.	1.5	244
45	Common variants at 19p13 are associated with susceptibility to ovarian cancer. <i>Nature Genetics</i> , 2010, 42, 880-884.	9.4	235
46	Genome-wide association study identifies three new melanoma susceptibility loci. <i>Nature Genetics</i> , 2011, 43, 1108-1113.	9.4	230
47	Mutational spectrum in a worldwide study of 29,700 families with <i>BRCA1</i> or <i>BRCA2</i> mutations. <i>Human Mutation</i> , 2018, 39, 593-620.	1.1	224
48	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	9.4	221
49	Reproductive risk factors for ovarian cancer in carriers of <i>BRCA1</i> or <i>BRCA2</i> mutations: a case-control study. <i>Lancet Oncology</i> , The, 2007, 8, 26-34.	5.1	220
50	Genome-wide association study of renal cell carcinoma identifies two susceptibility loci on 2p21 and 11q13.3. <i>Nature Genetics</i> , 2011, 43, 60-65.	9.4	220
51	Cancer prevention with aspirin in hereditary colorectal cancer (Lynch syndrome), 10-year follow-up and registry-based 20-year data in the CAPP2 study: a double-blind, randomised, placebo-controlled trial. <i>Lancet</i> , The, 2020, 395, 1855-1863.	6.3	220
52	Genome-wide meta-analysis identifies five new susceptibility loci for cutaneous malignant melanoma. <i>Nature Genetics</i> , 2015, 47, 987-995.	9.4	218
53	<i>RAD51</i> 135G→C Modifies Breast Cancer Risk among <i>BRCA2</i> Mutation Carriers: Results from a Combined Analysis of 19 Studies. <i>American Journal of Human Genetics</i> , 2007, 81, 1186-1200.	2.6	217
54	Risk of Breast Cancer in Women With a <i>CHEK2</i> Mutation With and Without a Family History of Breast Cancer. <i>Journal of Clinical Oncology</i> , 2011, 29, 3747-3752.	0.8	207

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55	Functional Variants at the 11q13 Risk Locus for Breast Cancer Regulate Cyclin D1 Expression through Long-Range Enhancers. <i>American Journal of Human Genetics</i> , 2013, 92, 489-503.	2.6	201
56	Integrated proteogenomic deep sequencing and analytics accurately identify non-canonical peptides in tumor immunopeptidomes. <i>Nature Communications</i> , 2020, 11, 1293.	5.8	196
57	Targeted Prostate Cancer Screening in BRCA1 and BRCA2 Mutation Carriers: Results from the Initial Screening Round of the IMPACT Study. <i>European Urology</i> , 2014, 66, 489-499.	0.9	195
58	Proteogenomic insights into the biology and treatment of HPV-negative head and neck squamous cell carcinoma. <i>Cancer Cell</i> , 2021, 39, 361-379.e16.	7.7	189
59	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
60	Hormone Therapy and the Risk of Breast Cancer in BRCA1 Mutation Carriers. <i>Journal of the National Cancer Institute</i> , 2008, 100, 1361-1367.	3.0	179
61	Results of a phase II open-label, non-randomized trial of cisplatin chemotherapy in patients with BRCA1-positive metastatic breast cancer. <i>Breast Cancer Research</i> , 2012, 14, R110.	2.2	179
62	<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
63	Molecular diagnosis of pituitary adenoma predisposition caused by aryl hydrocarbon receptor-interacting protein gene mutations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 4101-4105.	3.3	173
64	A high proportion of founder BRCA1 mutations in Polish breast cancer families. <i>International Journal of Cancer</i> , 2004, 110, 683-686.	2.3	170
65	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
66	Germline RECQL mutations are associated with breast cancer susceptibility. <i>Nature Genetics</i> , 2015, 47, 643-646.	9.4	168
67	Predictors of Contralateral Prophylactic Mastectomy in Women With a <i>BRCA1</i> or <i>BRCA2</i> Mutation: The Hereditary Breast Cancer Clinical Study Group. <i>Journal of Clinical Oncology</i> , 2008, 26, 1093-1097.	0.8	161
68	Bilateral Oophorectomy and Breast Cancer Risk in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	3.0	160
69	A Genome-Wide Association Study of Upper Aerodigestive Tract Cancers Conducted within the INHANCE Consortium. <i>PLoS Genetics</i> , 2011, 7, e1001333.	1.5	158
70	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
71	Effect of pregnancy as a risk factor for breast cancer in BRCA1/BRCA2 mutation carriers. <i>International Journal of Cancer</i> , 2005, 117, 988-991.	2.3	152
72	Low penetrance breast cancer susceptibility loci are associated with specific breast tumor subtypes: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2011, 20, 3289-3303.	1.4	152

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73	Age- and Tumor Subtype-Specific Breast Cancer Risk Estimates for <i>CH</i> and <i>EK</i> <i>2</i> *1100delC Carriers. <i>Journal of Clinical Oncology</i> , 2016, 34, 2750-2760.	0.8	152
74	Interim Results from the IMPACT Study: Evidence for Prostate-specific Antigen Screening in BRCA2 Mutation Carriers. <i>European Urology</i> , 2019, 76, 831-842.	0.9	148
75	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. <i>Nature Communications</i> , 2013, 4, 1628.	5.8	144
76	Genome-wide association meta-analyses combining multiple risk phenotypes provide insights into the genetic architecture of cutaneous melanoma susceptibility. <i>Nature Genetics</i> , 2020, 52, 494-504.	9.4	138
77	A Novel Founder CHEK2 Mutation is Associated with Increased Prostate Cancer Risk: Table 1. <i>Cancer Research</i> , 2004, 64, 2677-2679.	0.4	137
78	Clinical outcomes in women with breast cancer and a PALB2 mutation: a prospective cohort analysis. <i>Lancet Oncology</i> , The, 2015, 16, 638-644.	5.1	137
79	The risk of endometrial cancer in women with BRCA1 and BRCA2 mutations. A prospective study. <i>Gynecologic Oncology</i> , 2007, 104, 7-10.	0.6	135
80	Reproductive and Hormonal Factors, and Ovarian Cancer Risk for <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers: Results from the International <i>BRCA1/2</i> Carrier Cohort Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 601-610.	1.1	130
81	Breast cancer risk variants at 6q25 display different phenotype associations and regulate <i>ESR1</i> , <i>RMND1</i> and <i>CCDC170</i> . <i>Nature Genetics</i> , 2016, 48, 374-386.	9.4	125
82	Mutational Signature Analysis Reveals <i>NTHL1</i> Deficiency to Cause a Multi-tumor Phenotype. <i>Cancer Cell</i> , 2019, 35, 256-266.e5.	7.7	123
83	Hormone Replacement Therapy After Oophorectomy and Breast Cancer Risk Among <i>BRCA1</i> Mutation Carriers. <i>JAMA Oncology</i> , 2018, 4, 1059.	3.4	121
84	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
85	Estimation of Genetic Correlation via Linkage Disequilibrium Score Regression and Genomic Restricted Maximum Likelihood. <i>American Journal of Human Genetics</i> , 2018, 102, 1185-1194.	2.6	119
86	A meta-analysis of genome-wide association studies to identify prostate cancer susceptibility loci associated with aggressive and non-aggressive disease. <i>Human Molecular Genetics</i> , 2013, 22, 408-415.	1.4	118
87	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
88	Ten-Year Survival in Patients With <i>BRCA1</i> -Negative and <i>BRCA1</i> -Positive Breast Cancer. <i>Journal of Clinical Oncology</i> , 2013, 31, 3191-3196.	0.8	112
89	A Range of Cancers Is Associated with the rs6983267 Marker on Chromosome 8. <i>Cancer Research</i> , 2008, 68, 9982-9986.	0.4	111
90	A variant in <i>FTO</i> shows association with melanoma risk not due to BMI. <i>Nature Genetics</i> , 2013, 45, 428-432.	9.4	111

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91	Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016, 45, 1619-1630.	0.9	111
92	The Effect on Melanoma Risk of Genes Previously Associated With Telomere Length. <i>Journal of the National Cancer Institute</i> , 2014, 106, .	3.0	109
93	On the origin and diffusion of BRCA1 c.5266dupC (5382insC) in European populations. <i>European Journal of Human Genetics</i> , 2011, 19, 300-306.	1.4	107
94	The incidence of endometrial cancer in women with BRCA1 and BRCA2 mutations: An international prospective cohort study. <i>Gynecologic Oncology</i> , 2013, 130, 127-131.	0.6	106
95	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , 2017, 8, 15724.	5.8	106
96	The NOD2 3020insC Mutation and the Risk of Colorectal Cancer: Table 1. <i>Cancer Research</i> , 2004, 64, 1604-1606.	0.4	105
97	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	5.8	105
98	Screening mammography and risk of breast cancer in BRCA1 and BRCA2 mutation carriers: a case-control study. <i>Lancet Oncology</i> , The, 2006, 7, 402-406.	5.1	104
99	Germline deletions in the EPCAM gene as a cause of Lynch syndrome – literature review. <i>Hereditary Cancer in Clinical Practice</i> , 2013, 11, 9.	0.6	104
100	Changes in body weight and the risk of breast cancer in BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research</i> , 2005, 7, R833-43.	2.2	103
101	Prevalence of BRCA1 and BRCA2 germline mutations in patients with triple-negative breast cancer. <i>Breast Cancer Research and Treatment</i> , 2015, 150, 71-80.	1.1	103
102	A deletion in CHEK2 of 5,395bp predisposes to breast cancer in Poland. <i>Breast Cancer Research and Treatment</i> , 2007, 102, 119-122.	1.1	102
103	PARP-1 expression in breast cancer including BRCA1-associated, triple negative and basal-like tumors: possible implications for PARP-1 inhibitor therapy. <i>Breast Cancer Research and Treatment</i> , 2011, 127, 861-869.	1.1	102
104	International trends in the uptake of cancer risk reduction strategies in women with a BRCA1 or BRCA2 mutation. <i>British Journal of Cancer</i> , 2019, 121, 15-21.	2.9	101
105	Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with TERT expression. <i>Human Molecular Genetics</i> , 2013, 22, 2520-2528.	1.4	100
106	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
107	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> , 2013, 93, 1046-1060.	2.6	98
108	Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31. <i>Nature Communications</i> , 2013, 4, 1627.	5.8	98

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109	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> , 2014, 16, 3419.	2.2	97
110	CHEK2 mutations and the risk of papillary thyroid cancer. <i>International Journal of Cancer</i> , 2015, 137, 548-552.	2.3	97
111	Long-term effect of resistant starch on cancer risk in carriers of hereditary colorectal cancer: an analysis from the CAPP2 randomised controlled trial. <i>Lancet Oncology</i> , The, 2012, 13, 1242-1249.	5.1	95
112	No evidence that protein truncating variants in BRIP1 are associated with breast cancer risk: implications for gene panel testing. <i>Journal of Medical Genetics</i> , 2016, 53, 298-309.	1.5	94
113	Suspected hereditary nonpolyposis colorectal cancer. <i>Diseases of the Colon and Rectum</i> , 1999, 42, 710-715.	0.7	93
114	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	5.8	93
115	Breastfeeding and the risk of breast cancer in BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research</i> , 2012, 14, R42.	2.2	92
116	Cancer variation associated with the position of the mutation in the BRCA2 gene. <i>Familial Cancer</i> , 2002, 3, 1-10.	0.9	91
117	Interplay between BRCA1 and RHAMM Regulates Epithelial Apicobasal Polarization and May Influence Risk of Breast Cancer. <i>PLoS Biology</i> , 2011, 9, e1001199.	2.6	91
118	MiRNA-362a-3p induces cell cycle arrest through targeting of E2F1, USF2 and PTPN1 and is associated with recurrence of colorectal cancer. <i>International Journal of Cancer</i> , 2013, 133, 67-78.	2.3	91
119	Obesity, Aspirin, and Risk of Colorectal Cancer in Carriers of Hereditary Colorectal Cancer: A Prospective Investigation in the CAPP2 Study. <i>Journal of Clinical Oncology</i> , 2015, 33, 3591-3597.	0.8	91
120	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
121	Novel mutations in the LKB1/STK11 gene in Dutch Peutz-Jeghers families. <i>Human Mutation</i> , 1999, 13, 476-481.	1.1	89
122	Family History of Cancer and Cancer Risks in Women with BRCA1 or BRCA2 Mutations. <i>Journal of the National Cancer Institute</i> , 2010, 102, 1874-1878.	3.0	89
123	Male breast cancer in BRCA1 and BRCA2 mutation carriers: pathology data from the Consortium of Investigators of Modifiers of BRCA1/2. <i>Breast Cancer Research</i> , 2016, 18, 15.	2.2	88
124	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. <i>Nature Communications</i> , 2018, 9, 2256.	5.8	88
125	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88
126	Infertility, treatment of infertility, and the risk of breast cancer among women with BRCA1 and BRCA2 mutations: a case-control study. <i>Cancer Causes and Control</i> , 2008, 19, 1111-1119.	0.8	87

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127	Hereditary ovarian cancer in Poland. <i>International Journal of Cancer</i> , 2003, 106, 942-945.	2.3	82
128	Influence of selected lifestyle factors on breast and ovarian cancer risk in BRCA1 mutation carriers from Poland. <i>Breast Cancer Research and Treatment</i> , 2006, 95, 105-109.	1.1	82
129	Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of BRCA1 and BRCA2 pathogenic variants. <i>Genetics in Medicine</i> , 2020, 22, 1653-1666.	1.1	82
130	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. <i>Human Molecular Genetics</i> , 2012, 21, 456-462.	1.4	81
131	Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2019, 48, 795-806.	0.9	81
132	Germline 657del5 mutation in the NBS1 gene in breast cancer patients. <i>International Journal of Cancer</i> , 2003, 106, 379-381.	2.3	80
133	<i>Aryl hydrocarbon receptor interacting protein</i> (<i>AIP</i>) gene mutation analysis in children and adolescents with sporadic pituitary adenomas. <i>Clinical Endocrinology</i> , 2008, 69, 621-627.	1.2	80
134	Increased Incidence of Visceral Metastases in Scottish Patients With BRCA1/2-Defective Ovarian Cancer: An Extension of the Ovarian BRCAness Phenotype. <i>Journal of Clinical Oncology</i> , 2010, 28, 2505-2511.	0.8	80
135	Polymorphism in the P-glycoprotein drug transporter MDR1 gene in colon cancer patients. <i>European Journal of Clinical Pharmacology</i> , 2005, 61, 389-394.	0.8	79
136	Suspected HNPCC and Amsterdam criteria II: evaluation of mutation detection rate, an international collaborative study. <i>International Journal of Colorectal Disease</i> , 2002, 17, 109-114.	1.0	78
137	Common variants at 12p11, 12q24, 9p21, 9q31.2 and in ZNF365 are associated with breast cancer risk for BRCA1 and/or BRCA2 mutation carriers. <i>Breast Cancer Research</i> , 2012, 14, R33.	2.2	78
138	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	5.8	78
139	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. <i>Journal of the National Cancer Institute</i> , 2016, 108, djv315.	3.0	77
140	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
141	Genetic testing and common disorders in a public health framework: how to assess relevance and possibilities. <i>European Journal of Human Genetics</i> , 2011, 19, S6-S44.	1.4	75
142	<i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017, 77, 2789-2799.	0.4	75
143	Coffee consumption and breast cancer risk among BRCA1 and BRCA2 mutation carriers. <i>International Journal of Cancer</i> , 2006, 118, 103-107.	2.3	73
144	Age at menarche and the risk of breast cancer in BRCA1 and BRCA2 mutation carriers. <i>Cancer Causes and Control</i> , 2005, 16, 667-674.	0.8	71

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