

Jolanta Lissowska

List of Publications by Year in descending order

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Version: 2024-02-01

434
papers

49,291
citations

2427

97
h-index

2280

200
g-index

450
all docs

450
docs citations

450
times ranked

49602
citing authors

#	ARTICLE	IF	CITATIONS
1	Interleukin-1 polymorphisms associated with increased risk of gastric cancer. Nature, 2000, 404, 398-402.	27.8	2,197
2	Genome-wide association study identifies novel breast cancer susceptibility loci. Nature, 2007, 447, 1087-1093.	27.8	2,165
3	Genome-wide association study identifies 30 loci associated with bipolar disorder. Nature Genetics, 2019, 51, 793-803.	21.4	1,191
4	A susceptibility locus for lung cancer maps to nicotinic acetylcholine receptor subunit genes on 15q25. Nature, 2008, 452, 633-637.	27.8	1,169
5	Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94.	27.8	1,099
6	Analysis of shared heritability in common disorders of the brain. Science, 2018, 360, .	12.6	1,085
7	Genome-wide meta-analyses identify multiple loci associated with smoking behavior. Nature Genetics, 2010, 42, 441-447.	21.4	1,083
8	Human Papillomavirus and Oral Cancer: The International Agency for Research on Cancer Multicenter Study. Journal of the National Cancer Institute, 2003, 95, 1772-1783.	6.3	1,013
9	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. Nature Genetics, 2013, 45, 353-361.	21.4	960
10	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. Cell, 2019, 179, 1469-1482.e11.	28.9	935
11	Subtyping of Breast Cancer by Immunohistochemistry to Investigate a Relationship between Subtype and Short and Long Term Survival: A Collaborative Analysis of Data for 10,159 Cases from 12 Studies. PLoS Medicine, 2010, 7, e1000279.	8.4	764
12	Menarche, menopause, and breast cancer risk: individual participant meta-analysis, including 118,964 women with breast cancer from 117 epidemiological studies. Lancet Oncology, The, 2012, 13, 1141-1151.	10.7	753
13	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. American Journal of Human Genetics, 2019, 104, 21-34.	6.2	711
14	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. Nature Genetics, 2019, 51, 1207-1214.	21.4	641
15	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. Nature Genetics, 2021, 53, 817-829.	21.4	629
16	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. Cell, 2018, 173, 1705-1715.e16.	28.9	623
17	Type I and II Endometrial Cancers: Have They Different Risk Factors?. Journal of Clinical Oncology, 2013, 31, 2607-2618.	1.6	613
18	Associations of Breast Cancer Risk Factors With Tumor Subtypes: A Pooled Analysis From the Breast Cancer Association Consortium Studies. Journal of the National Cancer Institute, 2011, 103, 250-263.	6.3	596

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19	A common coding variant in CASP8 is associated with breast cancer risk. Nature Genetics, 2007, 39, 352-358.	21.4	591
20	Detectable clonal mosaicism and its relationship to aging and cancer. Nature Genetics, 2012, 44, 651-658.	21.4	519
21	Lung cancer susceptibility locus at 5p15.33. Nature Genetics, 2008, 40, 1404-1406.	21.4	514
22	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. Nature Genetics, 2013, 45, 371-384.	21.4	493
23	A multistage genome-wide association study in breast cancer identifies two new risk alleles at 1p11.2 and 14q24.1 (RAD51L1). Nature Genetics, 2009, 41, 579-584.	21.4	487
24	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. Nature Genetics, 2017, 49, 1126-1132.	21.4	472
25	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. Nature Genetics, 2009, 41, 585-590.	21.4	434
26	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. Journal of the National Cancer Institute, 2015, 107, .	6.3	428
27	Significant Locus and Metabolic Genetic Correlations Revealed in Genome-Wide Association Study of Anorexia Nervosa. American Journal of Psychiatry, 2017, 174, 850-858.	7.2	410
28	Differences in Risk Factors for Breast Cancer Molecular Subtypes in a Population-Based Study. Cancer Epidemiology Biomarkers and Prevention, 2007, 16, 439-443.	2.5	394
29	Cigarette smoking and lung cancer—relative risk estimates for the major histological types from a pooled analysis of case-control studies. International Journal of Cancer, 2012, 131, 1210-1219.	5.1	390
30	Performance of Common Genetic Variants in Breast-Cancer Risk Models. New England Journal of Medicine, 2010, 362, 986-993.	27.0	376
31	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. Nature Genetics, 2013, 45, 392-398.	21.4	374
32	Rare variants of large effect in BRCA2 and CHEK2 affect risk of lung cancer. Nature Genetics, 2014, 46, 736-741.	21.4	360
33	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. Nature Genetics, 2017, 49, 680-691.	21.4	356
34	Menopausal hormone use and ovarian cancer risk: individual participant meta-analysis of 52 epidemiological studies. Lancet, The, 2015, 385, 1835-1842.	13.7	349
35	Hormone-receptor expression and ovarian cancer survival: an Ovarian Tumor Tissue Analysis consortium study. Lancet Oncology, The, 2013, 14, 853-862.	10.7	335
36	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. Nature Genetics, 2013, 45, 362-370.	21.4	326

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37	A genome-wide association study identifies susceptibility loci for ovarian cancer at 2q31 and 8q24. <i>Nature Genetics</i> , 2010, 42, 874-879.	21.4	321
38	Oral Health and Risk of Squamous Cell Carcinoma of the Head and Neck and Esophagus: Results of Two Multicentric Case-Control Studies. <i>American Journal of Epidemiology</i> , 2007, 166, 1159-1173.	3.4	318
39	Heterogeneity of Breast Cancer Associations with Five Susceptibility Loci by Clinical and Pathological Characteristics. <i>PLoS Genetics</i> , 2008, 4, e1000054.	3.5	315
40	Improved survival of gastric cancer with tumour Epstein-Barr virus positivity: an international pooled analysis. <i>Gut</i> , 2014, 63, 236-243.	12.1	309
41	Genome-wide association study reveals two new risk loci for bipolar disorder. <i>Nature Communications</i> , 2014, 5, 3339.	12.8	294
42	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	21.4	289
43	A genome-wide association study of anorexia nervosa. <i>Molecular Psychiatry</i> , 2014, 19, 1085-1094.	7.9	282
44	A genome-wide association study identifies a new ovarian cancer susceptibility locus on 9p22.2. <i>Nature Genetics</i> , 2009, 41, 996-1000.	21.4	276
45	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.	21.4	265
46	Genome-wide association analysis identifies three new breast cancer susceptibility loci. <i>Nature Genetics</i> , 2012, 44, 312-318.	21.4	256
47	A Functional Polymorphism of Toll-Like Receptor 4 Gene Increases Risk of Gastric Carcinoma and Its Precursors. <i>Gastroenterology</i> , 2007, 132, 905-912.	1.3	247
48	Common variants at 19p13 are associated with susceptibility to ovarian cancer. <i>Nature Genetics</i> , 2010, 42, 880-884.	21.4	235
49	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	21.4	221
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.	21.4	220
51	Cigarette, Cigar, and Pipe Smoking and the Risk of Head and Neck Cancers: Pooled Analysis in the International Head and Neck Cancer Epidemiology Consortium. <i>American Journal of Epidemiology</i> , 2013, 178, 679-690.	3.4	220
52	Genetic identification of cell types underlying brain complex traits yields insights into the etiology of Parkinson's disease. <i>Nature Genetics</i> , 2020, 52, 482-493.	21.4	216
53	Cessation of alcohol drinking, tobacco smoking and the reversal of head and neck cancer risk. <i>International Journal of Epidemiology</i> , 2010, 39, 182-196.	1.9	210
54	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.	6.2	201

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55	Influence of common genetic variation on lung cancer risk: meta-analysis of 14 900 cases and 29 485 controls. <i>Human Molecular Genetics</i> , 2012, 21, 4980-4995.	2.9	196
56	Predictors of global methylation levels in blood DNA of healthy subjects: a combined analysis. <i>International Journal of Epidemiology</i> , 2012, 41, 126-139.	1.9	187
57	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. <i>American Journal of Psychiatry</i> , 2019, 176, 651-660.	7.2	186
58	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.	21.4	184
59	Correction: The role of interleukin-1 polymorphisms in the pathogenesis of gastric cancer. <i>Nature</i> , 2001, 412, 99-99.	27.8	183
60	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018, 9, 3166.	12.8	178
61	<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.	3.2	174
62	Obesity and risk of ovarian cancer subtypes: evidence from the Ovarian Cancer Association Consortium. <i>Endocrine-Related Cancer</i> , 2013, 20, 251-262.	3.1	169
63	Large-Scale Investigation of Base Excision Repair Genetic Polymorphisms and Lung Cancer Risk in a Multicenter Study. <i>Journal of the National Cancer Institute</i> , 2005, 97, 567-576.	6.3	166
64	Low human papillomavirus prevalence in head and neck cancer: results from two large case-control studies in high-incidence regions. <i>International Journal of Epidemiology</i> , 2011, 40, 489-502.	1.9	165
65	Genome-wide association analyses identify new susceptibility loci for oral cavity and pharyngeal cancer. <i>Nature Genetics</i> , 2016, 48, 1544-1550.	21.4	164
66	Multiple ADH genes are associated with upper aerodigestive cancers. <i>Nature Genetics</i> , 2008, 40, 707-709.	21.4	161
67	Previous Lung Diseases and Lung Cancer Risk: A Pooled Analysis From the International Lung Cancer Consortium. <i>American Journal of Epidemiology</i> , 2012, 176, 573-585.	3.4	160
68	A Genome-Wide Association Study of Upper Aerodigestive Tract Cancers Conducted within the INHANCE Consortium. <i>PLoS Genetics</i> , 2011, 7, e1001333.	3.5	158
69	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.	2.9	152
70	Smoking, alcohol, diet, dentition and sexual practices in the epidemiology of oral cancer in Poland. <i>European Journal of Cancer Prevention</i> , 2003, 12, 25-33.	1.3	151
71	Exposure to Diesel Motor Exhaust and Lung Cancer Risk in a Pooled Analysis from Case-Control Studies in Europe and Canada. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2011, 183, 941-948.	5.6	150
72	Genome-wide association study of borderline personality disorder reveals genetic overlap with bipolar disorder, major depression and schizophrenia. <i>Translational Psychiatry</i> , 2017, 7, e1155-e1155.	4.8	150

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73	Investigation of the fine structure of European populations with applications to disease association studies. <i>European Journal of Human Genetics</i> , 2008, 16, 1413-1429.	2.8	147
74	Polymorphisms in DNA double-strand break repair genes and risk of breast cancer: two population-based studies in USA and Poland, and meta-analyses. <i>Human Genetics</i> , 2006, 119, 376-388.	3.8	144
75	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. <i>Nature Communications</i> , 2013, 4, 1628.	12.8	144
76	Total Exposure and Exposure Rate Effects for Alcohol and Smoking and Risk of Head and Neck Cancer: A Pooled Analysis of Case-Control Studies. <i>American Journal of Epidemiology</i> , 2009, 170, 937-947.	3.4	143
77	Increased risk of lung cancer in individuals with a family history of the disease: A pooled analysis from the International Lung Cancer Consortium. <i>European Journal of Cancer</i> , 2012, 48, 1957-1968.	2.8	143
78	The Genetics of the Mood Disorder Spectrum: Genome-wide Association Analyses of More Than 185,000 Cases and 439,000 Controls. <i>Biological Psychiatry</i> , 2020, 88, 169-184.	1.3	137
79	Effect of cruciferous vegetables on lung cancer in patients stratified by genetic status: a mendelian randomisation approach. <i>Lancet, The</i> , 2005, 366, 1558-1560.	13.7	136
80	Evidence of Gene-Environment Interactions between Common Breast Cancer Susceptibility Loci and Established Environmental Risk Factors. <i>PLoS Genetics</i> , 2013, 9, e1003284.	3.5	136
81	Improving genetic prediction by leveraging genetic correlations among human diseases and traits. <i>Nature Communications</i> , 2018, 9, 989.	12.8	136
82	Genetic Polymorphisms in Base-Excision Repair Pathway Genes and Risk of Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 353-358.	2.5	132
83	Risk factors for head and neck cancer in young adults: a pooled analysis in the INHANCE consortium. <i>International Journal of Epidemiology</i> , 2015, 44, 169-185.	1.9	128
84	Established breast cancer risk factors by clinically important tumour characteristics. <i>British Journal of Cancer</i> , 2006, 95, 123-129.	6.4	127
85	Association between a 15q25 gene variant, smoking quantity and tobacco-related cancers among 17 000 individuals. <i>International Journal of Epidemiology</i> , 2010, 39, 563-577.	1.9	125
86	Family history of cancer: Pooled analysis in the International Head and Neck Cancer Epidemiology Consortium. <i>International Journal of Cancer</i> , 2009, 124, 394-401.	5.1	122
87	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2022, 91, 313-327.	1.3	114
88	In-Home Coal and Wood Use and Lung Cancer Risk: A Pooled Analysis of the International Lung Cancer Consortium. <i>Environmental Health Perspectives</i> , 2010, 118, 1743-1747.	6.0	112
89	Estimating and explaining the effect of education and income on head and neck cancer risk: INHANCE consortium pooled analysis of 31 case-control studies from 27 countries. <i>International Journal of Cancer</i> , 2015, 136, 1125-1139.	5.1	112
90	Obesity and survival among women with ovarian cancer: results from the Ovarian Cancer Association Consortium. <i>British Journal of Cancer</i> , 2015, 113, 817-826.	6.4	111

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91	Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. International Journal of Epidemiology, 2016, 45, 1619-1630.	1.9	111
92	Lung Cancer and Indoor Pollution from Heating and Cooking with Solid Fuels. American Journal of Epidemiology, 2005, 162, 326-333.	3.4	110
93	Genetic polymorphisms in the one-carbon metabolism pathway and breast cancer risk: A population-based case-control study and meta-analyses. International Journal of Cancer, 2007, 120, 2696-2703.	5.1	107
94	Lung cancer and socioeconomic status in a pooled analysis of case-control studies. PLoS ONE, 2018, 13, e0192999.	2.5	107
95	Telomere Length in Peripheral Leukocyte DNA and Gastric Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 3103-3109.	2.5	106
96	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. Nature Communications, 2017, 8, 15724.	12.8	106
97	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. Nature Communications, 2014, 5, 4999.	12.8	105
98	Blood leukocyte DNA hypomethylation and gastric cancer risk in a high-risk Polish population. International Journal of Cancer, 2010, 127, 1866-1874.	5.1	103
99	Common Breast Cancer Susceptibility Variants in <i>LSP1</i> and <i>RAD51L1</i> Are Associated with Mammographic Density Measures that Predict Breast Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 1156-1166.	2.5	101
100	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. American Journal of Human Genetics, 2015, 96, 487-497.	6.2	101
101	The role of oral hygiene in head and neck cancer: results from International Head and Neck Cancer Epidemiology (INHANCE) consortium. Annals of Oncology, 2016, 27, 1619-1625.	1.2	101
102	19p13.1 Is a Triple-Negative-Specific Breast Cancer Susceptibility Locus. Cancer Research, 2012, 72, 1795-1803.	0.9	100
103	Exposure to secondhand tobacco smoke and lung cancer by histological type: A pooled analysis of the International Lung Cancer Consortium (ILCCO). International Journal of Cancer, 2014, 135, 1918-1930.	5.1	100
104	Risk of Estrogen Receptor-Positive and -Negative Breast Cancer and Single-Nucleotide Polymorphism 2q35-rs13387042. Journal of the National Cancer Institute, 2009, 101, 1012-1018.	6.3	99
105	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. Journal of the National Cancer Institute, 2015, 107, djv219.	6.3	99
106	Polymorphisms in Th1-type cell-mediated response genes and risk of gastric cancer. Carcinogenesis, 2007, 28, 118-123.	2.8	98
107	Fine-Scale Mapping of the FGFR2 Breast Cancer Risk Locus: Putative Functional Variants Differentially Bind FOXA1 and E2F1. American Journal of Human Genetics, 2013, 93, 1046-1060.	6.2	98
108	Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31. Nature Communications, 2013, 4, 1627.	12.8	98

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109	Is Previous Respiratory Disease a Risk Factor for Lung Cancer?. American Journal of Respiratory and Critical Care Medicine, 2014, 190, 549-559.	5.6	97
110	Refined histopathological predictors of BRCA1 and BRCA2 mutation status: a large-scale analysis of breast cancer characteristics from the BCAC, CIMBA, and ENIGMA consortia. Breast Cancer Research, 2014, 16, 3419.	5.0	97
111	Obesity and cancer: Mendelian randomization approach utilizing the FTO genotype. International Journal of Epidemiology, 2009, 38, 971-975.	1.9	96
112	Occupational Exposure to Crystalline Silica and Risk of Lung Cancer. Epidemiology, 2007, 18, 36-43.	2.7	94
113	International Lung Cancer Consortium: Pooled Analysis of Sequence Variants in DNA Repair and Cell Cycle Pathways. Cancer Epidemiology Biomarkers and Prevention, 2008, 17, 3081-3089.	2.5	93
114	Tobacco, alcohol, and diet in the etiology of laryngeal cancer: a population-based case-control study. Cancer Causes and Control, 1991, 2, 3-10.	1.8	92
115	DNA Repair and Cell Cycle Control Genes and the Risk of Young-Onset Lung Cancer. Cancer Research, 2006, 66, 11062-11069.	0.9	91
116	Risk of stomach cancer in relation to consumption of cigarettes, alcohol, tea and coffee in Warsaw, Poland. , 1999, 81, 871-876.		90
117	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. Human Molecular Genetics, 2014, 23, 6616-6633.	2.9	90
118	Body mass index and risk of head and neck cancer in a pooled analysis of case-control studies in the International Head and Neck Cancer Epidemiology (INHANCE) Consortium. International Journal of Epidemiology, 2010, 39, 1091-1102.	1.9	89
119	Development of lung cancer before the age of 50: the role of xenobiotic metabolizing genes. Carcinogenesis, 2007, 28, 1287-1293.	2.8	87
120	Occupational exposure to polycyclic aromatic hydrocarbons and lung cancer risk: a multicenter study in Europe. Occupational and Environmental Medicine, 2010, 67, 98-103.	2.8	86
121	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. Nature Communications, 2016, 7, 11843.	12.8	86
122	Prognostic gene expression signature for high-grade serous ovarian cancer. Annals of Oncology, 2020, 31, 1240-1250.	1.2	85
123	Association of ESR1 gene tagging SNPs with breast cancer risk. Human Molecular Genetics, 2009, 18, 1131-1139.	2.9	84
124	Cigarette smoking and risk of ovarian cancer: a pooled analysis of 21 case-control studies. Cancer Causes and Control, 2013, 24, 989-1004.	1.8	84
125	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. Breast Cancer Research, 2010, 12, R110.	5.0	82
126	The <sc>INHANCE</sc> consortium: toward a better understanding of the causes and mechanisms of head and neck cancer. Oral Diseases, 2015, 21, 685-693.	3.0	82

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127	Family history and lung cancer risk: international multicentre caseâ€“control study in Eastern and Central Europe and meta-analyses. <i>Cancer Causes and Control</i> , 2010, 21, 1091-1104.	1.8	81
128	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.	2.9	81
129	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.	1.9	81
130	The role of genetic breast cancer susceptibility variants as prognostic factors. <i>Human Molecular Genetics</i> , 2012, 21, 3926-3939.	2.9	80
131	Diet and Stomach Cancer Risk in Warsaw, Poland. <i>Nutrition and Cancer</i> , 2004, 48, 149-159.	2.0	79
132	Obesity, metabolic factors and risk of different histological types of lung cancer: A Mendelian randomization study. <i>PLoS ONE</i> , 2017, 12, e0177875.	2.5	79
133	Tagging Single Nucleotide Polymorphisms in Cell Cycle Control Genes and Susceptibility to Invasive Epithelial Ovarian Cancer. <i>Cancer Research</i> , 2007, 67, 3027-3035.	0.9	78
134	Genome-wide significant risk associations for mucinous ovarian carcinoma. <i>Nature Genetics</i> , 2015, 47, 888-897.	21.4	78
135	Association of Polygenic Liabilities for Major Depression, Bipolar Disorder, and Schizophrenia With Risk for Depression in the Danish Population. <i>JAMA Psychiatry</i> , 2019, 76, 516.	11.0	78
136	Identification of shared risk loci and pathways for bipolar disorder and schizophrenia. <i>PLoS ONE</i> , 2017, 12, e0171595.	2.5	77
137	A comprehensive analysis of common genetic variation in MUC1, MUC5AC, MUC6 genes and risk of stomach cancer. <i>Cancer Causes and Control</i> , 2010, 21, 313-321.	1.8	76
138	Age at Last Birth in Relation to Risk of Endometrial Cancer: Pooled Analysis in the Epidemiology of Endometrial Cancer Consortium. <i>American Journal of Epidemiology</i> , 2012, 176, 269-278.	3.4	76
139	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.	6.2	76
140	<i>ESR1/SYNE1</i> Polymorphism and Invasive Epithelial Ovarian Cancer Risk: An Ovarian Cancer Association Consortium Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 245-250.	2.5	75
141	Consortium analysis of 7 candidate SNPs for ovarian cancer. <i>International Journal of Cancer</i> , 2008, 123, 380-388.	5.1	73
142	Genetic variation in C20orf54, PLCE1 and MUC1 and the risk of upper gastrointestinal cancers in Caucasian populations. <i>European Journal of Cancer Prevention</i> , 2012, 21, 541-544.	1.3	72
143	The etiology of uterine sarcomas: a pooled analysis of the epidemiology of endometrial cancer consortium. <i>British Journal of Cancer</i> , 2013, 108, 727-734.	6.4	72
144	Occupational Exposure to Vinyl Chloride, Acrylonitrile and Styrene and Lung Cancer Risk (Europe). <i>Cancer Causes and Control</i> , 2004, 15, 445-452.	1.8	71

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145	Associations of common variants at 1p11.2 and 14q24.1 (RAD51L1) with breast cancer risk and heterogeneity by tumor subtype: findings from the Breast Cancer Association Consortium. Human Molecular Genetics, 2011, 20, 4693-4706.	2.9	71
146	Adult body mass index and risk of ovarian cancer by subtype: a Mendelian randomization study. International Journal of Epidemiology, 2016, 45, 884-895.	1.9	71
147	Exposure-Response Analyses of Asbestos and Lung Cancer Subtypes in a Pooled Analysis of Case-Control Studies. Epidemiology, 2017, 28, 288-299.	2.7	71
148	Genetic variation in five genes important in telomere biology and risk for breast cancer. British Journal of Cancer, 2007, 97, 832-836.	6.4	70
149	Genetic Variation in the Prostate Stem Cell Antigen Gene and Upper Gastrointestinal Cancer in White Individuals. Gastroenterology, 2011, 140, 435-441.	1.3	70
150	Asthma and lung cancer risk: a systematic investigation by the International Lung Cancer Consortium. Carcinogenesis, 2012, 33, 587-597.	2.8	69
151	Shared genetics underlying epidemiological association between endometriosis and ovarian cancer. Human Molecular Genetics, 2015, 24, 5955-5964.	2.9	68
152	High Cumulative Risk of Lung Cancer Death among Smokers and Nonsmokers in Central and Eastern Europe. American Journal of Epidemiology, 2006, 164, 1233-1241.	3.4	67
153	Genome-wide analysis implicates microRNAs and their target genes in the development of bipolar disorder. Translational Psychiatry, 2015, 5, e678-e678.	4.8	67
154	Uncommon CHEK2 mis-sense variant and reduced risk of tobacco-related cancers: case-control study. Human Molecular Genetics, 2007, 16, 1794-1801.	2.9	66
155	Adult height and head and neck cancer: a pooled analysis within the INHANCE Consortium. European Journal of Epidemiology, 2014, 29, 35-48.	5.7	66
156	Selected DNA repair polymorphisms and gastric cancer in Poland. Carcinogenesis, 2005, 26, 1354-1359.	2.8	65
157	Single Nucleotide Polymorphisms in the TP53 Region and Susceptibility to Invasive Epithelial Ovarian Cancer. Cancer Research, 2009, 69, 2349-2357.	0.9	63
158	Body Mass Index, Cigarette Smoking, and Alcohol Consumption and Cancers of the Oral Cavity, Pharynx, and Larynx: Modeling Odds Ratios in Pooled Case-Control Data. American Journal of Epidemiology, 2010, 171, 1250-1261.	3.4	63
159	Cis-eQTL analysis and functional validation of candidate susceptibility genes for high-grade serous ovarian cancer. Nature Communications, 2015, 6, 8234.	12.8	63
160	Evidence for three genetic loci involved in both anorexia nervosa risk and variation of body mass index. Molecular Psychiatry, 2017, 22, 192-201.	7.9	63
161	Genetic variation in tumor necrosis factor and lymphotoxin-alpha (TNF-LTA) and breast cancer risk. Human Genetics, 2007, 121, 483-490.	3.8	62
162	Discovery and validation of methylation markers for endometrial cancer. International Journal of Cancer, 2014, 135, 1860-1868.	5.1	62

#	ARTICLE	IF	CITATIONS
163	Two Estrogen-Related Variants in <i>CYP19A1</i> and Endometrial Cancer Risk: A Pooled Analysis in the Epidemiology of Endometrial Cancer Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 242-247.	2.5	61
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297	Common Genetic Variation in Circadian Rhythm Genes and Risk of Epithelial Ovarian Cancer (EOC). <i>Journal of Genetics and Genome Research</i> , 2015, 2, .	0.3	25
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303	Genetic variation in CYP17 and endometrial cancer risk. <i>Human Genetics</i> , 2008, 123, 155-162.	3.8	23
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305	Occupation and renal cell cancer in Central and Eastern Europe. <i>Occupational and Environmental Medicine</i> , 2010, 67, 47-53.	2.8	23
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