

Jolanta Lissowska

List of Publications by Year in descending order

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Version: 2024-02-01

434
papers

49,291
citations

2832

97
h-index

2688

199
g-index

450
all docs

450
docs citations

450
times ranked

53886
citing authors

#	ARTICLE	IF	CITATIONS
1	Interleukin-1 polymorphisms associated with increased risk of gastric cancer. <i>Nature</i> , 2000, 404, 398-402.	13.7	2,197
2	Genome-wide association study identifies novel breast cancer susceptibility loci. <i>Nature</i> , 2007, 447, 1087-1093.	13.7	2,165
3	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019, 51, 793-803.	9.4	1,191
4	A susceptibility locus for lung cancer maps to nicotinic acetylcholine receptor subunit genes on 15q25. <i>Nature</i> , 2008, 452, 633-637.	13.7	1,169
5	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099
6	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , 2018, 360, .	6.0	1,085
7	Genome-wide meta-analyses identify multiple loci associated with smoking behavior. <i>Nature Genetics</i> , 2010, 42, 441-447.	9.4	1,083
8	Human Papillomavirus and Oral Cancer: The International Agency for Research on Cancer Multicenter Study. <i>Journal of the National Cancer Institute</i> , 2003, 95, 1772-1783.	3.0	1,013
9	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	9.4	960
10	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , 2019, 179, 1469-1482.e11.	13.5	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. <i>PLoS Medicine</i> , 2010, 7, e1000279.	3.9	764
12	Menarche, menopause, and breast cancer risk: individual participant meta-analysis, including 118,964 women with breast cancer from 117 epidemiological studies. <i>Lancet Oncology</i> , The, 2012, 13, 1141-1151.	5.1	753
13	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
14	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. <i>Nature Genetics</i> , 2019, 51, 1207-1214.	9.4	641
15	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. <i>Nature Genetics</i> , 2021, 53, 817-829.	9.4	629
16	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. <i>Cell</i> , 2018, 173, 1705-1715.e16.	13.5	623
17	Type I and II Endometrial Cancers: Have They Different Risk Factors?. <i>Journal of Clinical Oncology</i> , 2013, 31, 2607-2618.	0.8	613
18	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

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19	A common coding variant in CASP8 is associated with breast cancer risk. <i>Nature Genetics</i> , 2007, 39, 352-358.	9.4	591
20	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , 2012, 44, 651-658.	9.4	519
21	Lung cancer susceptibility locus at 5p15.33. <i>Nature Genetics</i> , 2008, 40, 1404-1406.	9.4	514
22	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
23	A multistage genome-wide association study in breast cancer identifies two new risk alleles at 1p11.2 and 14q24.1 (RAD51L1). <i>Nature Genetics</i> , 2009, 41, 579-584.	9.4	487
24	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. <i>Nature Genetics</i> , 2017, 49, 1126-1132.	9.4	472
25	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. <i>Nature Genetics</i> , 2009, 41, 585-590.	9.4	434
26	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
27	Significant Locus and Metabolic Genetic Correlations Revealed in Genome-Wide Association Study of Anorexia Nervosa. <i>American Journal of Psychiatry</i> , 2017, 174, 850-858.	4.0	410
28	Differences in Risk Factors for Breast Cancer Molecular Subtypes in a Population-Based Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 439-443.	1.1	394
29	Cigarette smoking and lung cancer—relative risk estimates for the major histological types from a pooled analysis of case-control studies. <i>International Journal of Cancer</i> , 2012, 131, 1210-1219.	2.3	390
30	Performance of Common Genetic Variants in Breast-Cancer Risk Models. <i>New England Journal of Medicine</i> , 2010, 362, 986-993.	13.9	376
31	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	9.4	374
32	Rare variants of large effect in BRCA2 and CHEK2 affect risk of lung cancer. <i>Nature Genetics</i> , 2014, 46, 736-741.	9.4	360
33	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691.	9.4	356
34	Menopausal hormone use and ovarian cancer risk: individual participant meta-analysis of 52 epidemiological studies. <i>Lancet</i> , The, 2015, 385, 1835-1842.	6.3	349
35	Hormone-receptor expression and ovarian cancer survival: an Ovarian Tumor Tissue Analysis consortium study. <i>Lancet Oncology</i> , The, 2013, 14, 853-862.	5.1	335
36	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 362-370.	9.4	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.	9.4	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.	1.6	318
39	Heterogeneity of Breast Cancer Associations with Five Susceptibility Loci by Clinical and Pathological Characteristics. <i>PLoS Genetics</i> , 2008, 4, e1000054.	1.5	315
40	Improved survival of gastric cancer with tumour Epstein-Barr virus positivity: an international pooled analysis. <i>Gut</i> , 2014, 63, 236-243.	6.1	309
41	Genome-wide association study reveals two new risk loci for bipolar disorder. <i>Nature Communications</i> , 2014, 5, 3339.	5.8	294
42	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
43	A genome-wide association study of anorexia nervosa. <i>Molecular Psychiatry</i> , 2014, 19, 1085-1094.	4.1	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.	9.4	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.	9.4	265
46	Genome-wide association analysis identifies three new breast cancer susceptibility loci. <i>Nature Genetics</i> , 2012, 44, 312-318.	9.4	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.	0.6	247
48	Common variants at 19p13 are associated with susceptibility to ovarian cancer. <i>Nature Genetics</i> , 2010, 42, 880-884.	9.4	235
49	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	9.4	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.	9.4	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.	1.6	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.	9.4	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.	0.9	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.	2.6	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.	1.4	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.	0.9	187
57	CWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. <i>American Journal of Psychiatry</i> , 2019, 176, 651-660.	4.0	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.	9.4	184
59	Correction: The role of interleukin-1 polymorphisms in the pathogenesis of gastric cancer. <i>Nature</i> , 2001, 412, 99-99.	13.7	183
60	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018, 9, 3166.	5.8	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.	1.5	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.	1.6	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.	3.0	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.	0.9	165
65	Genome-wide association analyses identify new susceptibility loci for oral cavity and pharyngeal cancer. <i>Nature Genetics</i> , 2016, 48, 1544-1550.	9.4	164
66	Multiple ADH genes are associated with upper aerodigestive cancers. <i>Nature Genetics</i> , 2008, 40, 707-709.	9.4	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.	1.6	160
68	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
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.	1.4	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.	0.6	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.	2.5	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.	2.4	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.	1.4	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.	1.8	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.	5.8	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.	1.6	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.	1.3	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.	0.7	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.	6.3	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.	1.5	136
81	Improving genetic prediction by leveraging genetic correlations among human diseases and traits. <i>Nature Communications</i> , 2018, 9, 989.	5.8	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.	1.1	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.	0.9	128
84	Established breast cancer risk factors by clinically important tumour characteristics. <i>British Journal of Cancer</i> , 2006, 95, 123-129.	2.9	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.	0.9	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.	2.3	122
87	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2022, 91, 313-327.	0.7	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.	2.8	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.	2.3	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.	2.9	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	Lung Cancer and Indoor Pollution from Heating and Cooking with Solid Fuels. <i>American Journal of Epidemiology</i> , 2005, 162, 326-333.	1.6	110
93	Genetic polymorphisms in the one-carbon metabolism pathway and breast cancer risk: A population-based case-control study and meta-analyses. <i>International Journal of Cancer</i> , 2007, 120, 2696-2703.	2.3	107
94	Lung cancer and socioeconomic status in a pooled analysis of case-control studies. <i>PLoS ONE</i> , 2018, 13, e0192999.	1.1	107
95	Telomere Length in Peripheral Leukocyte DNA and Gastric Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 3103-3109.	1.1	106
96	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , 2017, 8, 15724.	5.8	106
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	Blood leukocyte DNA hypomethylation and gastric cancer risk in a high-risk Polish population. <i>International Journal of Cancer</i> , 2010, 127, 1866-1874.	2.3	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. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1156-1166.	1.1	101
100	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	2.6	101
101	The role of oral hygiene in head and neck cancer: results from International Head and Neck Cancer Epidemiology (INHANCE) consortium. <i>Annals of Oncology</i> , 2016, 27, 1619-1625.	0.6	101
102	19p13.1 Is a Triple-Negative-Specific Breast Cancer Susceptibility Locus. <i>Cancer Research</i> , 2012, 72, 1795-1803.	0.4	100
103	Exposure to secondhand tobacco smoke and lung cancer by histological type: A pooled analysis of the International Lung Cancer Consortium (ILCCO). <i>International Journal of Cancer</i> , 2014, 135, 1918-1930.	2.3	100
104	Risk of Estrogen Receptor-Positive and -Negative Breast Cancer and Single-Nucleotide Polymorphism 2q35-rs13387042. <i>Journal of the National Cancer Institute</i> , 2009, 101, 1012-1018.	3.0	99
105	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
106	Polymorphisms in Th1-type cell-mediated response genes and risk of gastric cancer. <i>Carcinogenesis</i> , 2007, 28, 118-123.	1.3	98
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	Is Previous Respiratory Disease a Risk Factor for Lung Cancer?. American Journal of Respiratory and Critical Care Medicine, 2014, 190, 549-559.	2.5	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.	2.2	97
111	Obesity and cancer: Mendelian randomization approach utilizing the FTO genotype. International Journal of Epidemiology, 2009, 38, 971-975.	0.9	96
112	Occupational Exposure to Crystalline Silica and Risk of Lung Cancer. Epidemiology, 2007, 18, 36-43.	1.2	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.	1.1	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.	0.8	92
115	DNA Repair and Cell Cycle Control Genes and the Risk of Young-Onset Lung Cancer. Cancer Research, 2006, 66, 11062-11069.	0.4	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.	1.4	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.	0.9	89
119	Development of lung cancer before the age of 50: the role of xenobiotic metabolizing genes. Carcinogenesis, 2007, 28, 1287-1293.	1.3	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.	1.3	86
121	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. Nature Communications, 2016, 7, 11843.	5.8	86
122	Prognostic gene expression signature for high-grade serous ovarian cancer. Annals of Oncology, 2020, 31, 1240-1250.	0.6	85
123	Association of ESR1 gene tagging SNPs with breast cancer risk. Human Molecular Genetics, 2009, 18, 1131-1139.	1.4	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.	0.8	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.	2.2	82
126	The INHANCE consortium: toward a better understanding of the causes and mechanisms of head and neck cancer. Oral Diseases, 2015, 21, 685-693.	1.5	82

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127	Family history and lung cancer risk: international multicentre case-control study in Eastern and Central Europe and meta-analyses. <i>Cancer Causes and Control</i> , 2010, 21, 1091-1104.	0.8	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.	1.4	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.	0.9	81
130	The role of genetic breast cancer susceptibility variants as prognostic factors. <i>Human Molecular Genetics</i> , 2012, 21, 3926-3939.	1.4	80
131	Diet and Stomach Cancer Risk in Warsaw, Poland. <i>Nutrition and Cancer</i> , 2004, 48, 149-159.	0.9	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.	1.1	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.	0.4	78
134	Genome-wide significant risk associations for mucinous ovarian carcinoma. <i>Nature Genetics</i> , 2015, 47, 888-897.	9.4	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.	6.0	78
136	Identification of shared risk loci and pathways for bipolar disorder and schizophrenia. <i>PLoS ONE</i> , 2017, 12, e0171595.	1.1	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.	0.8	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.	1.6	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.	2.6	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.	1.1	75
141	Consortium analysis of 7 candidate SNPs for ovarian cancer. <i>International Journal of Cancer</i> , 2008, 123, 380-388.	2.3	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.	0.6	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.	2.9	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.	0.8	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. <i>Human Molecular Genetics</i> , 2011, 20, 4693-4706.	1.4	71
146	Adult body mass index and risk of ovarian cancer by subtype: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016, 45, 884-895.	0.9	71
147	Exposure-Response Analyses of Asbestos and Lung Cancer Subtypes in a Pooled Analysis of Case-Control Studies. <i>Epidemiology</i> , 2017, 28, 288-299.	1.2	71
148	Genetic variation in five genes important in telomere biology and risk for breast cancer. <i>British Journal of Cancer</i> , 2007, 97, 832-836.	2.9	70
149	Genetic Variation in the Prostate Stem Cell Antigen Gene and Upper Gastrointestinal Cancer in White Individuals. <i>Gastroenterology</i> , 2011, 140, 435-441.	0.6	70
150	Asthma and lung cancer risk: a systematic investigation by the International Lung Cancer Consortium. <i>Carcinogenesis</i> , 2012, 33, 587-597.	1.3	69
151	Shared genetics underlying epidemiological association between endometriosis and ovarian cancer. <i>Human Molecular Genetics</i> , 2015, 24, 5955-5964.	1.4	68
152	High Cumulative Risk of Lung Cancer Death among Smokers and Nonsmokers in Central and Eastern Europe. <i>American Journal of Epidemiology</i> , 2006, 164, 1233-1241.	1.6	67
153	Genome-wide analysis implicates microRNAs and their target genes in the development of bipolar disorder. <i>Translational Psychiatry</i> , 2015, 5, e678-e678.	2.4	67
154	Uncommon CHEK2 mis-sense variant and reduced risk of tobacco-related cancers: case-control study. <i>Human Molecular Genetics</i> , 2007, 16, 1794-1801.	1.4	66
155	Adult height and head and neck cancer: a pooled analysis within the INHANCE Consortium. <i>European Journal of Epidemiology</i> , 2014, 29, 35-48.	2.5	66
156	Selected DNA repair polymorphisms and gastric cancer in Poland. <i>Carcinogenesis</i> , 2005, 26, 1354-1359.	1.3	65
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