

# Jolanta Lissowska

## List of Publications by Citations

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411  
papers

37,188  
citations

89  
h-index

180  
g-index

450  
ext. papers

44,000  
ext. citations

9.3  
avg, IF

6.42  
L-index

#	Paper	IF	Citations
411	Genome-wide association study identifies novel breast cancer susceptibility loci. <i>Nature</i> , <b>2007</b> , 447, 1087-93	50.4	1957
410	Interleukin-1 polymorphisms associated with increased risk of gastric cancer. <i>Nature</i> , <b>2000</b> , 404, 398-402	50.4	1924
409	A susceptibility locus for lung cancer maps to nicotinic acetylcholine receptor subunit genes on 15q25. <i>Nature</i> , <b>2008</b> , 452, 633-7	50.4	1003
408	Genome-wide meta-analyses identify multiple loci associated with smoking behavior. <i>Nature Genetics</i> , <b>2010</b> , 42, 441-7	36.3	927
407	Human papillomavirus and oral cancer: the International Agency for Research on Cancer multicenter study. <i>Journal of the National Cancer Institute</i> , <b>2003</b> , 95, 1772-83	9.7	864
406	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , <b>2013</b> , 45, 353-61, 361e1-2	36.3	813
405	Analysis of shared heritability in common disorders of the brain. <i>Science</i> , <b>2018</b> , 360,	33.3	666
404	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , <b>2019</b> , 51, 793-803	36.3	662
403	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , <b>2017</b> , 551, 92-94	50.4	643
402	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> , <b>2010</b> , 7, e1000279	11.6	616
401	A common coding variant in CASP8 is associated with breast cancer risk. <i>Nature Genetics</i> , <b>2007</b> , 39, 352-8	36.3	557
400	Menarche, menopause, and breast cancer risk: individual participant meta-analysis, including 118 964 women with breast cancer from 117 epidemiological studies. <i>Lancet Oncology, The</i> , <b>2012</b> , 13, 1141-51	21.7	529
399	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> , <b>2011</b> , 103, 250-63	9.7	513
398	Lung cancer susceptibility locus at 5p15.33. <i>Nature Genetics</i> , <b>2008</b> , 40, 1404-6	36.3	466
397	Type I and II endometrial cancers: have they different risk factors?. <i>Journal of Clinical Oncology</i> , <b>2013</b> , 31, 2607-18	2.2	458
396	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> , <b>2009</b> , 41, 579-84	36.3	452
395	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

394	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , <b>2012</b> , 44, 651-8	36.3	409
393	Genomic Relationships, Novel Loci, and Pleiotropic Mechanisms across Eight Psychiatric Disorders. <i>Cell</i> , <b>2019</b> , 179, 1469-1482.e11	56.2	402
392	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. <i>Nature Genetics</i> , <b>2009</b> , 41, 585-90	36.3	393
391	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
390	Genomic Dissection of Bipolar Disorder and Schizophrenia, Including 28 Subphenotypes. <i>Cell</i> , <b>2018</b> , 173, 1705-1715.e16	56.2	360
389	Differences in risk factors for breast cancer molecular subtypes in a population-based study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2007</b> , 16, 439-43	4	349
388	Performance of common genetic variants in breast-cancer risk models. <i>New England Journal of Medicine</i> , <b>2010</b> , 362, 986-93	59.2	334
387	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
386	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
385	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. <i>Nature Genetics</i> , <b>2019</b> , 51, 1207-1214	36.3	303
384	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
383	A genome-wide association study identifies susceptibility loci for ovarian cancer at 2q31 and 8q24. <i>Nature Genetics</i> , <b>2010</b> , 42, 874-9	36.3	277
382	Significant Locus and Metabolic Genetic Correlations Revealed in Genome-Wide Association Study of Anorexia Nervosa. <i>American Journal of Psychiatry</i> , <b>2017</b> , 174, 850-858	11.9	276
381	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> , <b>2012</b> , 131, 1210-9	7.5	275
380	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. <i>Nature Genetics</i> , <b>2013</b> , 45, 362-70, 370e1-2	36.3	267
379	Menopausal hormone use and ovarian cancer risk: individual participant meta-analysis of 52 epidemiological studies. <i>Lancet, The</i> , <b>2015</b> , 385, 1835-42	40	267
378	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> , <b>2007</b> , 166, 1159-73	3.8	261
377	Genome-wide association study reveals two new risk loci for bipolar disorder. <i>Nature Communications</i> , <b>2014</b> , 5, 3339	17.4	248

376	Hormone-receptor expression and ovarian cancer survival: an Ovarian Tumor Tissue Analysis consortium study. <i>Lancet Oncology, The</i> , <b>2013</b> , 14, 853-62	21.7	248
375	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. <i>Nature Genetics</i> , <b>2017</b> , 49, 1126-1132	36.3	246
374	A genome-wide association study identifies a new ovarian cancer susceptibility locus on 9p22.2. <i>Nature Genetics</i> , <b>2009</b> , 41, 996-1000	36.3	240
373	Improved survival of gastric cancer with tumour Epstein-Barr virus positivity: an international pooled analysis. <i>Gut</i> , <b>2014</b> , 63, 236-43	19.2	238
372	Genome-wide association analysis identifies three new breast cancer susceptibility loci. <i>Nature Genetics</i> , <b>2012</b> , 44, 312-8	36.3	237
371	Rare variants of large effect in BRCA2 and CHEK2 affect risk of lung cancer. <i>Nature Genetics</i> , <b>2014</b> , 46, 736-41	36.3	228
370	A functional polymorphism of toll-like receptor 4 gene increases risk of gastric carcinoma and its precursors. <i>Gastroenterology</i> , <b>2007</b> , 132, 905-12	13.3	225
369	A genome-wide association study of anorexia nervosa. <i>Molecular Psychiatry</i> , <b>2014</b> , 19, 1085-94	15.1	224
368	Common variants at 19p13 are associated with susceptibility to ovarian cancer. <i>Nature Genetics</i> , <b>2010</b> , 42, 880-4	36.3	210
367	Genome-wide association study of renal cell carcinoma identifies two susceptibility loci on 2p21 and 11q13.3. <i>Nature Genetics</i> , <b>2011</b> , 43, 60-5	36.3	199
366	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , <b>2017</b> , 49, 680-691	36.3	190
365	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
364	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , <b>2015</b> , 47, 164-71	36.3	177
363	Predictors of global methylation levels in blood DNA of healthy subjects: a combined analysis. <i>International Journal of Epidemiology</i> , <b>2012</b> , 41, 126-39	7.8	177
362	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> , <b>2013</b> , 92, 489-503	11	167
361	Cessation of alcohol drinking, tobacco smoking and the reversal of head and neck cancer risk. <i>International Journal of Epidemiology</i> , <b>2010</b> , 39, 182-96	7.8	163
360	Influence of common genetic variation on lung cancer risk: meta-analysis of 14 900 cases and 29 485 controls. <i>Human Molecular Genetics</i> , <b>2012</b> , 21, 4980-95	5.6	159
359	The role of interleukin-1 polymorphisms in the pathogenesis of gastric cancer. <i>Nature</i> , <b>2001</b> , 412, 99	50.4	157

358	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> , <b>2013</b> , 178, 679-90	3.8	152
357	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> , <b>2005</b> , 97, 567-76	9.7	152
356	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> , <b>2011</b> , 40, 489-502	7.8	148
355	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> , <b>2011</b> , 20, 3289-303	5.6	140
354	Multiple ADH genes are associated with upper aerodigestive cancers. <i>Nature Genetics</i> , <b>2008</b> , 40, 707-9	36.3	140
353	Smoking, alcohol, diet, dentition and sexual practices in the epidemiology of oral cancer in Poland. <i>European Journal of Cancer Prevention</i> , <b>2003</b> , 12, 25-33	2	139
352	Investigation of the fine structure of European populations with applications to disease association studies. <i>European Journal of Human Genetics</i> , <b>2008</b> , 16, 1413-29	5.3	137
351	A genome-wide association study of upper aerodigestive tract cancers conducted within the INHANCE consortium. <i>PLoS Genetics</i> , <b>2011</b> , 7, e1001333	6	136
350	Obesity and risk of ovarian cancer subtypes: evidence from the Ovarian Cancer Association Consortium. <i>Endocrine-Related Cancer</i> , <b>2013</b> , 20, 251-62	5.7	135
349	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> , <b>2006</b> , 119, 376-88	6.3	133
348	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. <i>Nature Communications</i> , <b>2013</b> , 4, 1628	17.4	124
347	Previous lung diseases and lung cancer risk: a pooled analysis from the International Lung Cancer Consortium. <i>American Journal of Epidemiology</i> , <b>2012</b> , 176, 573-85	3.8	123
346	Genetic polymorphisms in base-excision repair pathway genes and risk of breast cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2006</b> , 15, 353-8	4	122
345	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
344	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> , <b>2011</b> , 183, 941-8	10.2	116
343	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> , <b>2009</b> , 170, 937-47	3.8	114
342	Effect of cruciferous vegetables on lung cancer in patients stratified by genetic status: a mendelian randomisation approach. <i>Lancet, The</i> , <b>2005</b> , 366, 1558-60	40	114
341	Evidence of gene-environment interactions between common breast cancer susceptibility loci and established environmental risk factors. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003284	6	112

340	Established breast cancer risk factors by clinically important tumour characteristics. <i>British Journal of Cancer</i> , <b>2006</b> , 95, 123-9	8.7	112
339	Association between a 15q25 gene variant, smoking quantity and tobacco-related cancers among 17 000 individuals. <i>International Journal of Epidemiology</i> , <b>2010</b> , 39, 563-77	7.8	110
338	GWAS of Suicide Attempt in Psychiatric Disorders and Association With Major Depression Polygenic Risk Scores. <i>American Journal of Psychiatry</i> , <b>2019</b> , 176, 651-660	11.9	103
337	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> , <b>2007</b> , 120, 2696-703	7.5	102
336	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
335	Family history of cancer: pooled analysis in the International Head and Neck Cancer Epidemiology Consortium. <i>International Journal of Cancer</i> , <b>2009</b> , 124, 394-401	7.5	101
334	Genome-wide association study of borderline personality disorder reveals genetic overlap with bipolar disorder, major depression and schizophrenia. <i>Translational Psychiatry</i> , <b>2017</b> , 7, e1155	8.6	100
333	Risk factors for head and neck cancer in young adults: a pooled analysis in the INHANCE consortium. <i>International Journal of Epidemiology</i> , <b>2015</b> , 44, 169-85	7.8	98
332	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> , <b>2012</b> , 48, 1957-68	7.5	98
331	Telomere length in peripheral leukocyte DNA and gastric cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2009</b> , 18, 3103-9	4	96
330	Blood leukocyte DNA hypomethylation and gastric cancer risk in a high-risk Polish population. <i>International Journal of Cancer</i> , <b>2010</b> , 127, 1866-74	7.5	94
329	19p13.1 is a triple-negative-specific breast cancer susceptibility locus. <i>Cancer Research</i> , <b>2012</b> , 72, 1795-803	8.1	93
328	Lung cancer and indoor pollution from heating and cooking with solid fuels: the IARC international multicentre case-control study in Eastern/Central Europe and the United Kingdom. <i>American Journal of Epidemiology</i> , <b>2005</b> , 162, 326-33	3.8	93
327	Genome-wide association analyses identify new susceptibility loci for oral cavity and pharyngeal cancer. <i>Nature Genetics</i> , <b>2016</b> , 48, 1544-1550	36.3	92
326	Common breast cancer susceptibility variants in LSP1 and RAD51L1 are associated with mammographic density measures that predict breast cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2012</b> , 21, 1156-66	4	92
325	In-home coal and wood use and lung cancer risk: a pooled analysis of the International Lung Cancer Consortium. <i>Environmental Health Perspectives</i> , <b>2010</b> , 118, 1743-7	8.4	91
324	Risk of estrogen receptor-positive and -negative breast cancer and single-nucleotide polymorphism 2q35-rs13387042. <i>Journal of the National Cancer Institute</i> , <b>2009</b> , 101, 1012-8	9.7	90
323	Polymorphisms in Th1-type cell-mediated response genes and risk of gastric cancer. <i>Carcinogenesis</i> , <b>2007</b> , 28, 118-23	4.6	89

322	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
321	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> , <b>2015</b> , 136, 1125-39	7.5	86
320	International Lung Cancer Consortium: pooled analysis of sequence variants in DNA repair and cell cycle pathways. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2008</b> , 17, 3081-9	4	86
319	Identification and molecular characterization of a new ovarian cancer susceptibility locus at 17q21.31. <i>Nature Communications</i> , <b>2013</b> , 4, 1627	17.4	85
318	Risk of stomach cancer in relation to consumption of cigarettes, alcohol, tea and coffee in Warsaw, Poland. <i>International Journal of Cancer</i> , <b>1999</b> , 81, 871-6	7.5	85
317	Genome-wide association study of more than 40,000 bipolar disorder cases provides new insights into the underlying biology. <i>Nature Genetics</i> , <b>2021</b> , 53, 817-829	36.3	83
316	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
315	Obesity and cancer: Mendelian randomization approach utilizing the FTO genotype. <i>International Journal of Epidemiology</i> , <b>2009</b> , 38, 971-5	7.8	82
314	DNA repair and cell cycle control genes and the risk of young-onset lung cancer. <i>Cancer Research</i> , <b>2006</b> , 66, 11062-9	10.1	82
313	Development of lung cancer before the age of 50: the role of xenobiotic metabolizing genes. <i>Carcinogenesis</i> , <b>2007</b> , 28, 1287-93	4.6	82
312	Obesity and survival among women with ovarian cancer: results from the Ovarian Cancer Association Consortium. <i>British Journal of Cancer</i> , <b>2015</b> , 113, 817-26	8.7	80
311	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
310	Genetic identification of cell types underlying brain complex traits yields insights into the etiology of Parkinson disease. <i>Nature Genetics</i> , <b>2020</b> , 52, 482-493	36.3	79
309	Occupational exposure to crystalline silica and risk of lung cancer: a multicenter case-control study in Europe. <i>Epidemiology</i> , <b>2007</b> , 18, 36-43	3.1	79
308	Tobacco, alcohol, and diet in the etiology of laryngeal cancer: a population-based case-control study. <i>Cancer Causes and Control</i> , <b>1991</b> , 2, 3-10	2.8	78
307	Characterization of large structural genetic mosaicism in human autosomes. <i>American Journal of Human Genetics</i> , <b>2015</b> , 96, 487-97	11	77
306	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 6616-33	5.6	77
305	Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , <b>2016</b> , 45, 1619-1630	7.8	77

304	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. <i>Nature Genetics</i> , <b>2020</b> , 52, 572-581	36.3	76
303	Improving genetic prediction by leveraging genetic correlations among human diseases and traits. <i>Nature Communications</i> , <b>2018</b> , 9, 989	17.4	76
302	Association of ESR1 gene tagging SNPs with breast cancer risk. <i>Human Molecular Genetics</i> , <b>2009</b> , 18, 1131-9	3.9	75
301	The role of genetic breast cancer susceptibility variants as prognostic factors. <i>Human Molecular Genetics</i> , <b>2012</b> , 21, 3926-39	5.6	75
300	Tagging single nucleotide polymorphisms in cell cycle control genes and susceptibility to invasive epithelial ovarian cancer. <i>Cancer Research</i> , <b>2007</b> , 67, 3027-35	10.1	75
299	Diet and stomach cancer risk in Warsaw, Poland. <i>Nutrition and Cancer</i> , <b>2004</b> , 48, 149-59	2.8	75
298	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
297	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. <i>Human Molecular Genetics</i> , <b>2012</b> , 21, 456-62	5.6	74
296	Assessing interactions between the associations of common genetic susceptibility variants, reproductive history and body mass index with breast cancer risk in the breast cancer association consortium: a combined case-control study. <i>Breast Cancer Research</i> , <b>2010</b> , 12, R110	8.3	74
295	Is previous respiratory disease a risk factor for lung cancer?. <i>American Journal of Respiratory and Critical Care Medicine</i> , <b>2014</b> , 190, 549-59	10.2	73
294	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , <b>2018</b> , 9, 3166	17.4	70
293	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. <i>International Journal of Epidemiology</i> , <b>2010</b> , 39, 1091-102	7.8	70
292	Cigarette smoking and risk of ovarian cancer: a pooled analysis of 21 case-control studies. <i>Cancer Causes and Control</i> , <b>2013</b> , 24, 989-1004	2.8	69
291	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> , <b>2014</b> , 135, 1918-30	7.5	69
290	Occupational exposure to polycyclic aromatic hydrocarbons and lung cancer risk: a multicenter study in Europe. <i>Occupational and Environmental Medicine</i> , <b>2010</b> , 67, 98-103	2.1	68
289	A comprehensive analysis of common genetic variation in MUC1, MUC5AC, MUC6 genes and risk of stomach cancer. <i>Cancer Causes and Control</i> , <b>2010</b> , 21, 313-21	2.8	68
288	The INHANCE consortium: toward a better understanding of the causes and mechanisms of head and neck cancer. <i>Oral Diseases</i> , <b>2015</b> , 21, 685-93	3.5	66
287	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> , <b>2011</b> , 20, 4693-706	5.6	66



286	Consortium analysis of 7 candidate SNPs for ovarian cancer. <i>International Journal of Cancer</i> , <b>2008</b> , 123, 380-388	7.5	66
285	Genetic variation in the prostate stem cell antigen gene and upper gastrointestinal cancer in white individuals. <i>Gastroenterology</i> , <b>2011</b> , 140, 435-41	13.3	65
284	Genetic variation in five genes important in telomere biology and risk for breast cancer. <i>British Journal of Cancer</i> , <b>2007</b> , 97, 832-6	8.7	65
283	ESR1/SYNE1 polymorphism and invasive epithelial ovarian cancer risk: an Ovarian Cancer Association Consortium study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2010</b> , 19, 245-50	4	64
282	Genetic variation in C20orf54, PLCE1 and MUC1 and the risk of upper gastrointestinal cancers in Caucasian populations. <i>European Journal of Cancer Prevention</i> , <b>2012</b> , 21, 541-4	2	64
281	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> , <b>2016</b> , 27, 1619-25	10.3	62
280	Genome-wide significant risk associations for mucinous ovarian carcinoma. <i>Nature Genetics</i> , <b>2015</b> , 47, 888-97	36.3	60
279	The etiology of uterine sarcomas: a pooled analysis of the epidemiology of endometrial cancer consortium. <i>British Journal of Cancer</i> , <b>2013</b> , 108, 727-34	8.7	60
278	Genetic variation in tumor necrosis factor and lymphotoxin-alpha (TNF-LTA) and breast cancer risk. <i>Human Genetics</i> , <b>2007</b> , 121, 483-90	6.3	60
277	High cumulative risk of lung cancer death among smokers and nonsmokers in Central and Eastern Europe. <i>American Journal of Epidemiology</i> , <b>2006</b> , 164, 1233-41	3.8	60
276	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , <b>2016</b> , 7, 11843	17.4	59
275	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
274	Selected DNA repair polymorphisms and gastric cancer in Poland. <i>Carcinogenesis</i> , <b>2005</b> , 26, 1354-9	4.6	59
273	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> , <b>2020</b> , 88, 169-184	7.9	57
272	DNA hypermethylation of ESR1 and PGR in breast cancer: pathologic and epidemiologic associations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2009</b> , 18, 3036-43	4	56
271	Obesity, metabolic factors and risk of different histological types of lung cancer: A Mendelian randomization study. <i>PLoS ONE</i> , <b>2017</b> , 12, e0177875	3.7	56
270	Two estrogen-related variants in CYP19A1 and endometrial cancer risk: a pooled analysis in the Epidemiology of Endometrial Cancer Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2009</b> , 18, 242-7	4	55
269	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> , <b>2012</b> , 176, 269-78	3.8	55

268	Leukocyte telomere length in a population-based case-control study of ovarian cancer: a pilot study. <i>Cancer Causes and Control</i> , <b>2010</b> , 21, 77-82	2.8	55
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258	The ATM missense mutation p.Ser49Cys (c.146C>G) and the risk of breast cancer. <i>Human Mutation</i> , <b>2006</b> , 27, 538-44	4.7	52
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251	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> , <b>2010</b> , 21, 1091-104	2.8	50

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248	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
247	Relationship of mammographic density and gene expression: analysis of normal breast tissue surrounding breast cancer. <i>Clinical Cancer Research</i> , <b>2013</b> , 19, 4972-4982	12.9	48
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241	Association between a germline OCA2 polymorphism at chromosome 15q13.1 and estrogen receptor-negative breast cancer survival. <i>Journal of the National Cancer Institute</i> , <b>2010</b> , 102, 650-62	9.7	45
240	Expression of TGF-beta signaling factors in invasive breast cancers: relationships with age at diagnosis and tumor characteristics. <i>Breast Cancer Research and Treatment</i> , <b>2010</b> , 121, 727-35	4.4	45
239	Human papillomavirus infection in women with and without cervical cancer in Warsaw, Poland. <i>European Journal of Cancer</i> , <b>2008</b> , 44, 557-64	7.5	45
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