

# Stig Bojesen

## List of Publications by Year in descending order

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Version: 2024-02-01

294  
papers

30,638  
citations

5268

83  
h-index

5829

161  
g-index

326  
all docs

326  
docs citations

326  
times ranked

33258  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association study identifies novel breast cancer susceptibility loci. <i>Nature</i> , 2007, 447, 1087-1093.	27.8	2,165
2	Worldwide trends in hypertension prevalence and progress in treatment and control from 1990 to 2019: a pooled analysis of 1201 population-representative studies with 104 million participants. <i>Lancet</i> , The, 2021, 398, 957-980.	13.7	1,289
3	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	27.8	1,099
4	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	21.4	960
5	Statin Use and Reduced Cancer-Related Mortality. <i>New England Journal of Medicine</i> , 2012, 367, 1792-1802.	27.0	798
6	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019, 104, 21-34.	6.2	711
7	Statin Use and Reduced Cancer-Related Mortality. <i>New England Journal of Medicine</i> , 2013, 368, 574-577.	27.0	615
8	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.	6.3	596
9	A common coding variant in <i>CASP8</i> is associated with breast cancer risk. <i>Nature Genetics</i> , 2007, 39, 352-358.	21.4	591
10	Parent-of-origin-specific allelic associations among 106 genomic loci for age at menarche. <i>Nature</i> , 2014, 514, 92-97.	27.8	548
11	Breast Cancer Risk Genes " Association Analysis in More than 113,000 Women. <i>New England Journal of Medicine</i> , 2021, 384, 428-439.	27.0	532
12	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.	21.4	513
13	Multiple independent variants at the <i>TERT</i> locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384.	21.4	493
14	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. <i>Nature Genetics</i> , 2013, 45, 385-391.	21.4	492
15	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.	21.4	472
16	C-reactive Protein As a Predictor of Prognosis in Chronic Obstructive Pulmonary Disease. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2007, 175, 250-255.	5.6	456
17	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. <i>Nature Genetics</i> , 2009, 41, 585-590.	21.4	434
18	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	6.3	428

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19	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. <i>Nature Genetics</i> , 2017, 49, 834-841.	21.4	426
20	MicroRNA Biomarkers in Whole Blood for Detection of Pancreatic Cancer. <i>JAMA - Journal of the American Medical Association</i> , 2014, 311, 392.	7.4	380
21	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	21.4	374
22	Baseline C-Reactive Protein Is Associated With Incident Cancer and Survival in Patients With Cancer. <i>Journal of Clinical Oncology</i> , 2009, 27, 2217-2224.	1.6	359
23	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. <i>Nature Genetics</i> , 2015, 47, 1294-1303.	21.4	357
24	Heterogeneity of Breast Cancer Associations with Five Susceptibility Loci by Clinical and Pathological Characteristics. <i>PLoS Genetics</i> , 2008, 4, e1000054.	3.5	315
25	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	21.4	289
26	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 126-135.	2.5	278
27	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. <i>Nature Genetics</i> , 2011, 43, 785-791.	21.4	265
28	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
29	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.	21.4	264
30	<i> CHEK2</i> *1100delC Genotyping for Clinical Assessment of Breast Cancer Risk: Meta-Analyses of 26,000 Patient Cases and 27,000 Controls. <i>Journal of Clinical Oncology</i> , 2008, 26, 542-548.	1.6	262
31	Peripheral Blood Leukocyte Telomere Length and Mortality Among 64 637 Individuals From the General Population. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv074.	6.3	258
32	Genome-wide association analysis identifies three new breast cancer susceptibility loci. <i>Nature Genetics</i> , 2012, 44, 312-318.	21.4	256
33	Genetically low vitamin D concentrations and increased mortality: mendelian randomisation analysis in three large cohorts. <i>BMJ, The</i> , 2014, 349, g6330-g6330.	6.0	238
34	Low 25-Hydroxyvitamin D and Risk of Type 2 Diabetes: A Prospective Cohort Study and Metaanalysis. <i>Clinical Chemistry</i> , 2013, 59, 381-391.	3.2	236
35	Height and body-mass index trajectories of school-aged children and adolescents from 1985 to 2019 in 200 countries and territories: a pooled analysis of 2181 population-based studies with 65 million participants. <i>Lancet, The</i> , 2020, 396, 1511-1524.	13.7	219
36	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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37	Short Telomere Length, Cancer Survival, and Cancer Risk in 47102 Individuals. <i>Journal of the National Cancer Institute</i> , 2013, 105, 459-468.	6.3	195
38	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
39	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	27.8	183
40	<i>CHKB</i> , <i>CHKE2</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
41	Short Telomere Length, Myocardial Infarction, Ischemic Heart Disease, and Early Death. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2012, 32, 822-829.	2.4	172
42	Reduced 25-hydroxyvitamin D and risk of Alzheimer's disease and vascular dementia. <i>Alzheimer's and Dementia</i> , 2014, 10, 296-302.	0.8	164
43	<i>CHKE2</i> *1100delC Heterozygosity in Women With Breast Cancer Associated With Early Death, Breast Cancer-Specific Death, and Increased Risk of a Second Breast Cancer. <i>Journal of Clinical Oncology</i> , 2012, 30, 4308-4316.	1.6	162
44	Detection and characterization of lung cancer using cell-free DNA fragmentomes. <i>Nature Communications</i> , 2021, 12, 5060.	12.8	161
45	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.	9.4	157
46	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
47	Vitamin D concentration, obesity, and risk of diabetes: a mendelian randomisation study. <i>Lancet Diabetes and Endocrinology</i> , 2014, 2, 298-306.	11.4	152
48	Age- and Tumor Subtype-Specific Breast Cancer Risk Estimates for <i>CHKE2</i> *1100delC Carriers. <i>Journal of Clinical Oncology</i> , 2016, 34, 2750-2760.	1.6	152
49	<i>AHRH</i> (cg05575921) hypomethylation marks smoking behaviour, morbidity and mortality. <i>Thorax</i> , 2017, 72, 646-653.	5.6	147
50	Telomere Shortening Unrelated to Smoking, Body Weight, Physical Activity, and Alcohol Intake: 4,576 General Population Individuals with Repeat Measurements 10 Years Apart. <i>PLoS Genetics</i> , 2014, 10, e1004191.	3.5	139
51	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
52	Short telomere length, lung function and chronic obstructive pulmonary disease in 46,396 individuals. <i>Thorax</i> , 2013, 68, 429-435.	5.6	134
53	A genome-wide association scan (GWAS) for mean telomere length within the COGS project: identified loci show little association with hormone-related cancer risk. <i>Human Molecular Genetics</i> , 2013, 22, 5056-5064.	2.9	130
54	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.	21.4	125

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55	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	21.4	120
56	Lymphopenia and risk of infection and infection-related death in 98,344 individuals from a prospective Danish population-based study. <i>PLoS Medicine</i> , 2018, 15, e1002685.	8.4	119
57	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.	2.9	118
58	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.	8.4	118
59	Plasma YKL-40 levels in healthy subjects from the general population. <i>Clinica Chimica Acta</i> , 2011, 412, 709-712.	1.1	115
60	Tumor suppressor p53 Arg72Pro polymorphism and longevity, cancer survival, and risk of cancer in the general population. <i>Journal of Experimental Medicine</i> , 2007, 204, 1295-1301.	8.5	111
61	Increased Risk of Breast Cancer Associated With CHEK2*1100delC. <i>Journal of Clinical Oncology</i> , 2007, 26, 57-63.	1.6	110
62	The JAK2 V617F somatic mutation, mortality and cancer risk in the general population. <i>Haematologica</i> , 2011, 96, 450-453.	3.5	110
63	Serum Biomarker Signature-Based Liquid Biopsy for Diagnosis of Early-Stage Pancreatic Cancer. <i>Journal of Clinical Oncology</i> , 2018, 36, 2887-2894.	1.6	108
64	Combined genetic and splicing analysis of BRCA1 c.[594-2A>C; 641A>G] highlights the relevance of naturally occurring in-frame transcripts for developing disease gene variant classification algorithms. <i>Human Molecular Genetics</i> , 2016, 25, 2256-2268.	2.9	106
65	Identification of a BRCA2-Specific Modifier Locus at 6p24 Related to Breast Cancer Risk. <i>PLoS Genetics</i> , 2013, 9, e1003173.	3.5	105
66	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	12.8	105
67	Telomeres and human health. <i>Journal of Internal Medicine</i> , 2013, 274, 399-413.	6.0	104
68	C-Reactive Protein and the Risk of Cancer: A Mendelian Randomization Study. <i>Journal of the National Cancer Institute</i> , 2010, 102, 202-206.	6.3	103
69	19p13.1 Is a Triple-Negative-Specific Breast Cancer Susceptibility Locus. <i>Cancer Research</i> , 2012, 72, 1795-1803.	0.9	100
70	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.	2.9	100
71	Genome-wide association study identifies a common variant in RAD51B associated with male breast cancer risk. <i>Nature Genetics</i> , 2012, 44, 1182-1184.	21.4	99
72	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv219.	6.3	99

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73	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.	6.2	98
74	Smoking and Increased White and Red Blood Cells. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2019, 39, 965-977.	2.4	98
75	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.	5.0	97
76	Increased Risk for Other Cancers in Addition to Breast Cancer for CHEK2*1100delC Heterozygotes Estimated From the Copenhagen General Population Study. <i>Journal of Clinical Oncology</i> , 2016, 34, 1208-1216.	1.6	97
77	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.	3.2	94
78	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	12.8	93
79	Short Telomere Length and Ischemic Heart Disease: Observational and Genetic Studies in 290 022 Individuals. <i>Clinical Chemistry</i> , 2016, 62, 1140-1149.	3.2	93
80	Long telomeres and cancer risk among 95%568 individuals from the general population. <i>International Journal of Epidemiology</i> , 2016, 45, 1634-1643.	1.9	90
81	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	12.8	90
82	Association of Clinical Benign Prostate Hyperplasia with Prostate Cancer Incidence and Mortality Revisited: A Nationwide Cohort Study of 3 009 258 Men. <i>European Urology</i> , 2011, 60, 691-698.	1.9	89
83	Low Plasma 25-Hydroxyvitamin D and Risk of Tobacco-Related Cancer. <i>Clinical Chemistry</i> , 2013, 59, 771-780.	3.2	89
84	Risk of cancer among HIV-infected individuals compared to the background population. <i>Aids</i> , 2014, 28, 1499-1508.	2.2	89
85	Joint associations of a polygenic risk score and environmental risk factors for breast cancer in the Breast Cancer Association Consortium. <i>International Journal of Epidemiology</i> , 2018, 47, 526-536.	1.9	88
86	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	12.8	88
87	Diagnostic value of JAK2 V617F somatic mutation for myeloproliferative cancer in 49 488 individuals from the general population. <i>British Journal of Haematology</i> , 2013, 160, 70-79.	2.5	87
88	Prevalence of the HOXB13 G84E germline mutation in British men and correlation with prostate cancer risk, tumour characteristics and clinical outcomes. <i>Annals of Oncology</i> , 2015, 26, 756-761.	1.2	85
89	Association of ESR1 gene tagging SNPs with breast cancer risk. <i>Human Molecular Genetics</i> , 2009, 18, 1131-1139.	2.9	84
90	JAK2V617F somatic mutation in the general population: myeloproliferative neoplasm development and progression rate. <i>Haematologica</i> , 2014, 99, 1448-1455.	3.5	82

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91	Increased Body Mass Index, Elevated C-reactive Protein, and Short Telomere Length. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2014, 99, E1671-E1675.	3.6	81
92	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
93	The common germline Arg72Pro polymorphism of p53 and increased longevity in humans. <i>Cell Cycle</i> , 2008, 7, 158-163.	2.6	80
94	The role of genetic breast cancer susceptibility variants as prognostic factors. <i>Human Molecular Genetics</i> , 2012, 21, 3926-3939.	2.9	80
95	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
96	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	12.8	78
97	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.	6.3	77
98	Elevated Plasma YKL-40 Predicts Increased Risk of Gastrointestinal Cancer and Decreased Survival After Any Cancer Diagnosis in the General Population. <i>Journal of Clinical Oncology</i> , 2009, 27, 572-578.	1.6	76
99	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
100	BRCA2 Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017, 77, 2789-2799.	0.9	75
101	Plasma 25-hydroxyvitamin D, lung function and risk of chronic obstructive pulmonary disease. <i>Thorax</i> , 2014, 69, 24-31.	5.6	73
102	Inflammatory biomarkers and risk of cancer in 84,000 individuals from the general population. <i>International Journal of Cancer</i> , 2016, 139, 1493-1500.	5.1	73
103	Causal relationships between body mass index, smoking and lung cancer: Univariable and multivariable Mendelian randomization. <i>International Journal of Cancer</i> , 2021, 148, 1077-1086.	5.1	73
104	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.	2.9	71
105	Integrin $\beta$ 3 Leu33Pro Homozygosity and Risk of Cancer. <i>Journal of the National Cancer Institute</i> , 2003, 95, 1150-1157.	6.3	68
106	Elevated plasma YKL-40 levels and ischemic stroke in the general population. <i>Annals of Neurology</i> , 2010, 68, 672-680.	5.3	68
107	Multiple novel prostate cancer susceptibility signals identified by fine-mapping of known risk loci among Europeans. <i>Human Molecular Genetics</i> , 2015, 24, 5589-5602.	2.9	67
108	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017, 19, 599-603.	2.4	67

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109	Shorter leukocyte telomere length is associated with higher risk of infections: a prospective study of 75,309 individuals from the general population. <i>Haematologica</i> , 2017, 102, 1457-1465.	3.5	63
110	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2016, 23, 77-91.	3.1	62
111	Platelet glycoprotein IIb/IIIa PIA2/PIA2homozygosity associated with risk of ischemic cardiovascular disease and myocardial infarction in young men. <i>Journal of the American College of Cardiology</i> , 2003, 42, 661-667.	2.8	61
112	Identification of susceptibility pathways for the role of chromosome 15q25.1 in modifying lung cancer risk. <i>Nature Communications</i> , 2018, 9, 3221.	12.8	60
113	Evidence that the 5p12 Variant rs10941679 Confers Susceptibility to Estrogen-Receptor-Positive Breast Cancer through FGF10 and MRPS30 Regulation. <i>American Journal of Human Genetics</i> , 2016, 99, 903-911.	6.2	59
114	Is high vitamin B12 status a cause of lung cancer?. <i>International Journal of Cancer</i> , 2019, 145, 1499-1503.	5.1	58
115	Plasma YKL-40 and Total and Disease-Specific Mortality in the General Population. <i>Clinical Chemistry</i> , 2010, 56, 1580-1591.	3.2	57
116	Hyperhomocysteinemia, methylenetetrahydrofolate reductase c.677C>T polymorphism and risk of cancer: Cross-sectional and prospective studies and meta-analyses of 75,000 cases and 93,000 controls. <i>International Journal of Cancer</i> , 2011, 128, 644-652.	5.1	57
117	DNA mismatch repair gene MSH6 implicated in determining age at natural menopause. <i>Human Molecular Genetics</i> , 2014, 23, 2490-2497.	2.9	56
118	Identification of Novel Genetic Markers of Breast Cancer Survival. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	6.3	56
119	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2014, 23, 6096-6111.	2.9	53
120	Appraising the causal relevance of DNA methylation for risk of lung cancer. <i>International Journal of Epidemiology</i> , 2019, 48, 1493-1504.	1.9	53
121	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019, 120, 647-657.	6.4	52
122	Nicotinic Acetylcholine Receptor Polymorphism, Smoking Behavior, and Tobacco-Related Cancer and Lung and Cardiovascular Diseases: A Cohort Study. <i>Journal of Clinical Oncology</i> , 2011, 29, 2875-2882.	1.6	51
123	Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. <i>International Journal of Cancer</i> , 2016, 139, 1303-1317.	5.1	51
124	Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes. <i>JAMA Oncology</i> , 2022, 8, e216744.	7.1	51
125	Role of inflammatory marker YKL-40 in the diagnosis, prognosis and cause of cardiovascular and liver diseases. <i>Critical Reviews in Clinical Laboratory Sciences</i> , 2016, 53, 396-408.	6.1	50
126	Assessing Lung Cancer Absolute Risk Trajectory Based on a Polygenic Risk Model. <i>Cancer Research</i> , 2021, 81, 1607-1615.	0.9	50



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127	MicroRNA Related Polymorphisms and Breast Cancer Risk. PLoS ONE, 2014, 9, e109973.	2.5	49
128	Association Between a Germline OCA2 Polymorphism at Chromosome 15q13.1 and Estrogen Receptor- $\alpha$ Negative Breast Cancer Survival. Journal of the National Cancer Institute, 2010, 102, 650-662.	6.3	48
129	Plasma 25-Hydroxyvitamin D and Risk of Non-Melanoma and Melanoma Skin Cancer: A Prospective Cohort Study. Journal of Investigative Dermatology, 2013, 133, 629-636.	0.7	46
130	Arterial and venous thrombosis by high platelet count and high hematocrit: 108,521 individuals from the Copenhagen General Population Study. Journal of Thrombosis and Haemostasis, 2019, 17, 1898-1911.	3.8	46
131	Low high-density lipoprotein and increased risk of several cancers: 2 population-based cohort studies including 116,728 individuals. Journal of Hematology and Oncology, 2020, 13, 129.	17.0	46
132	Elevated Plasma YKL-40, Lipids and Lipoproteins, and Ischemic Vascular Disease in the General Population. Stroke, 2015, 46, 329-335.	2.0	45
133	Body mass index and breast cancer survival: a Mendelian randomization analysis. International Journal of Epidemiology, 2017, 46, 1814-1822.	1.9	45
134	Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. Journal of the National Cancer Institute, 2021, 113, 329-337.	6.3	45
135	<i>CHRNA3</i> genotype, nicotine dependence, lung function and disease in the general population. European Respiratory Journal, 2012, 40, 1538-1544.	6.7	44
136	Genetic predisposition to ductal carcinoma in situ of the breast. Breast Cancer Research, 2016, 18, 22.	5.0	43
137	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. Breast Cancer Research, 2017, 19, 119.	5.0	43
138	Fine mapping of MHC region in lung cancer highlights independent susceptibility loci by ethnicity. Nature Communications, 2018, 9, 3927.	12.8	43
139	Plasma testosterone in the general population, cancer prognosis and cancer risk: a prospective cohort study. Annals of Oncology, 2014, 25, 712-718.	1.2	42
140	High tobacco consumption is causally associated with increased all-cause mortality in a general population sample of 55,568 individuals, but not with short telomeres: a Mendelian randomization study. International Journal of Epidemiology, 2014, 43, 1473-1483.	1.9	41
141	Clinical value of serum hyaluronan and propeptide of type III collagen in patients with pancreatic cancer. International Journal of Cancer, 2020, 146, 2913-2922.	5.1	41
142	Heterogeneous contributions of change in population distribution of body mass index to change in obesity and underweight. ELife, 2021, 10, .	6.0	41
143	Prostate-Specific Antigen and Long-Term Prediction of Prostate Cancer Incidence and Mortality in the General Population. European Urology, 2012, 61, 865-874.	1.9	40
144	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. Human Molecular Genetics, 2015, 24, 2966-2984.	2.9	40

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145	Breast cancer risk factors and their effects on survival: a Mendelian randomisation study. <i>BMC Medicine</i> , 2020, 18, 327.	5.5	40
146	Genetic Predisposition to In Situ and Invasive Lobular Carcinoma of the Breast. <i>PLoS Genetics</i> , 2014, 10, e1004285.	3.5	39
147	Patient survival and tumor characteristics associated with CHEK2:p.I157T findings from the Breast Cancer Association Consortium. <i>Breast Cancer Research</i> , 2016, 18, 98.	5.0	39
148	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.	6.2	39
149	Secular trends in smoking in relation to prevalent and incident smoking-related disease: A prospective population-based study. <i>Tobacco Induced Diseases</i> , 2019, 17, 72.	0.6	39
150	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. <i>Human Molecular Genetics</i> , 2015, 24, 285-298.	2.9	38
151	Alcohol Consumption and Survival after a Breast Cancer Diagnosis: A Literature-Based Meta-analysis and Collaborative Analysis of Data for 29,239 Cases. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 934-945.	2.5	37
152	IgE and risk of cancer in 37 747 individuals from the general population. <i>Annals of Oncology</i> , 2015, 26, 1784-1790.	1.2	37
153	Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression. <i>American Journal of Human Genetics</i> , 2015, 97, 22-34.	6.2	37
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