

Stig Bojesen

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

292
papers

20,912
citations

73
h-index

139
g-index

326
ext. papers

25,866
ext. citations

10
avg, IF

6.29
L-index

#	Paper	IF	Citations
292	Rare germline copy number variants (CNVs) and breast cancer risk.. <i>Communications Biology</i> , 2022 , 5, 65	6.7	0
291	Genome-wide interaction analysis identified low-frequency variants with sex disparity in lung cancer risk.. <i>Human Molecular Genetics</i> , 2022 ,	5.6	1
290	Gene-gene interaction of AhRwith and within the Wntcascade affects susceptibility to lung cancer.. <i>European Journal of Medical Research</i> , 2022 , 27, 14	4.8	
289	Common variants in breast cancer risk loci predispose to distinct tumor subtypes.. <i>Breast Cancer Research</i> , 2022 , 24, 2	8.3	3
288	Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes.. <i>JAMA Oncology</i> , 2022 ,	13.4	4
287	Development and validation of a model to predict incident chronic liver disease in the general population: the CLiVD score.. <i>Journal of Hepatology</i> , 2022 ,	13.4	1
286	Self-reported and genetically predicted coffee consumption and smoking in dementia: A Mendelian randomization study.. <i>Atherosclerosis</i> , 2022 , 348, 36-43	3.1	1
285	Breast cancer risks associated with missense variants in breast cancer susceptibility genes.. <i>Genome Medicine</i> , 2022 , 14, 51	14.4	0
284	Allostatic load as predictor of mortality: a cohort study from Lolland-Falster, Denmark. <i>BMJ Open</i> , 2022 , 12, e057136	3	1
283	Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. <i>Scientific Reports</i> , 2021 , 11, 19787	4.9	0
282	Response to the letter entitled: Re: Pre-treatment serum vitamin D deficiency is associated with increased inflammatory biomarkers and short overall survival in patients with pancreatic cancer: Analysis of the prognostic effect of serum vitamin D on pancreatic cancer: Several confounders. <i>European Journal of Cancer</i> , 2021 , 156, 212-216	7.5	
281	Estimating dose-response relationships for vitamin D with coronary heart disease, stroke, and all-cause mortality: observational and Mendelian randomisation analyses. <i>Lancet Diabetes and Endocrinology</i> , 2021 , 9, 837-846	18.1	17
280	Circulating Protein Biomarkers for Use in Pancreatic Ductal Adenocarcinoma Identification. <i>Clinical Cancer Research</i> , 2021 , 27, 2592-2603	12.9	8
279	Heterogeneous contributions of change in population distribution of body mass index to change in obesity and underweight. <i>ELife</i> , 2021 , 10,	8.9	10
278	Genome-wide association meta-analysis identifies pleiotropic risk loci for aerodigestive squamous cell cancers. <i>PLoS Genetics</i> , 2021 , 17, e1009254	6	2
277	Genetic predisposition to long telomeres is associated with increased mortality after melanoma: A study of 2101 melanoma patients from hospital clinics and the general population. <i>Pigment Cell and Melanoma Research</i> , 2021 , 34, 946-954	4.5	0
276	Risk of ulcerative colitis and Crohn's disease in smokers lacks causal evidence. <i>European Journal of Epidemiology</i> , 2021 , 1	12.1	2

275	Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. <i>American Journal of Human Genetics</i> , 2021 , 108, 1190-1203	11	1
274	Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2021 , 113, 329-337	9.7	14
273	Common Susceptibility Loci for Male Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2021 , 113, 453-461	9.7	4
272	Evaluating the role of alcohol consumption in breast and ovarian cancer susceptibility using population-based cohort studies and two-sample Mendelian randomization analyses. <i>International Journal of Cancer</i> , 2021 , 148, 1338-1350	7.5	4
271	Integration of multiomic annotation data to prioritize and characterize inflammation and immune-related risk variants in squamous cell lung cancer. <i>Genetic Epidemiology</i> , 2021 , 45, 99-114	2.6	2
270	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	7.5	18
269	Myocardial Ischemia Induced by 5-Fluorouracil: A Prospective Electrocardiographic and Cardiac Biomarker Study. <i>Oncologist</i> , 2021 , 26, e403-e413	5.7	6
268	Comprehensive functional annotation of susceptibility variants identifies genetic heterogeneity between lung adenocarcinoma and squamous cell carcinoma. <i>Frontiers of Medicine</i> , 2021 , 15, 275-291	12	6
267	CYP3A7*1C allele: linking premenopausal oestrone and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , 2021 , 124, 842-854	8.7	2
266	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	36.3	62
265	Assessing Lung Cancer Absolute Risk Trajectory Based on a Polygenic Risk Model. <i>Cancer Research</i> , 2021 , 81, 1607-1615	10.1	6
264	Reference intervals for 12 clinical laboratory tests in a Danish population: The Lolland-Falster Health Study. <i>Scandinavian Journal of Clinical and Laboratory Investigation</i> , 2021 , 81, 104-111	2	0
263	A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. <i>Nature Communications</i> , 2021 , 12, 1078	17.4	4
262	Breast Cancer Risk Genes - Association Analysis in More than 113,000 Women. <i>New England Journal of Medicine</i> , 2021 , 384, 428-439	59.2	143
261	Pre-treatment serum vitamin D deficiency is associated with increased inflammatory biomarkers and short overall survival in patients with pancreatic cancer. <i>European Journal of Cancer</i> , 2021 , 144, 72-80	7.5	8
260	Association of germline genetic variants with breast cancer-specific survival in patient subgroups defined by clinic-pathological variables related to tumor biology and type of systemic treatment. <i>Breast Cancer Research</i> , 2021 , 23, 86	8.3	1
259	Mendelian randomisation study of smoking exposure in relation to breast cancer risk. <i>British Journal of Cancer</i> , 2021 , 125, 1135-1145	8.7	0
258	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021 , 596, 393-397	30.4	28

257	Detection and characterization of lung cancer using cell-free DNA fragmentomes. <i>Nature Communications</i> , 2021 , 12, 5060	17.4	21
256	Chronic lymphocytic leukaemia clones are detectable decades before diagnosis. <i>British Journal of Haematology</i> , 2021 ,	4.5	0
255	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, The</i> , 2021 , 398, 957-980	40	154
254	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021 , 30, 623-642	4	4
253	Pre- and Perioperative Inflammatory Biomarkers in Older Patients Resected for Localized Colorectal Cancer: Associations with Complications and Prognosis.. <i>Cancers</i> , 2021 , 14,	6.6	1
252	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	40	73
251	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	36.3	76
250	Protein-altering germline mutations implicate novel genes related to lung cancer development. <i>Nature Communications</i> , 2020 , 11, 2220	17.4	6
249	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , 2020 , 10, 9688	4.9	2
248	Reply to: Clinical impact of high platelet count and high hematocrit, by Marc Sorigue. <i>Journal of Thrombosis and Haemostasis</i> , 2020 , 18, 522-523	15.4	
247	Two-fold risk of pneumonia and respiratory mortality in individuals with myeloproliferative neoplasm: A population-based cohort study. <i>EClinicalMedicine</i> , 2020 , 21, 100295	11.3	2
246	Tocilizumab and soluble interleukin-6 receptor in V617F somatic mutation and myeloproliferative neoplasm. <i>EClinicalMedicine</i> , 2020 , 22, 100337	11.3	1
245	Transcriptome-wide association study of breast cancer risk by estrogen-receptor status. <i>Genetic Epidemiology</i> , 2020 , 44, 442-468	2.6	9
244	Incidental lymphopenia and mortality: a prospective cohort study. <i>Cmaj</i> , 2020 , 192, E25-E33	3.5	11
243	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020 , 11, 312	17.4	20
242	Elevated plasma YKL-40 and risk of infectious disease: a prospective study of 94665 individuals from the general population. <i>Clinical Microbiology and Infection</i> , 2020 , 26, 1411.e1-1411.e9	9.5	4
241	Prediction of contralateral breast cancer: external validation of risk calculators in 20 international cohorts. <i>Breast Cancer Research and Treatment</i> , 2020 , 181, 423-434	4.4	7
240	Transcriptome-wide association study reveals candidate causal genes for lung cancer. <i>International Journal of Cancer</i> , 2020 , 146, 1862-1878	7.5	8

239	Prognostic Value of Combined Detection of Serum IL6, YKL-40, and C-reactive Protein in Patients with Unresectable Pancreatic Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020 , 29, 176-184 ⁴		5
238	Genome-wide association study of INDELs identified four novel susceptibility loci associated with lung cancer risk. <i>International Journal of Cancer</i> , 2020 , 146, 2855-2864	7.5	2
237	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020 , 52, 56-73	36.3	56
236	Immune-mediated genetic pathways resulting in pulmonary function impairment increase lung cancer susceptibility. <i>Nature Communications</i> , 2020 , 11, 27	17.4	7
235	Smoking, blood cells and myeloproliferative neoplasms: meta-analysis and Mendelian randomization of 2B million people. <i>British Journal of Haematology</i> , 2020 , 189, 323-334	4.5	20
234	Low high-density lipoprotein and increased risk of several cancers: 2 population-based cohort studies including 116,728 individuals. <i>Journal of Hematology and Oncology</i> , 2020 , 13, 129	22.4	11
233	AHRR hypomethylation as an epigenetic marker of smoking history predicts risk of myocardial infarction in former smokers. <i>Atherosclerosis</i> , 2020 , 312, 8-15	3.1	2
232	Loss-of-function polymorphism in reduces risk of V617F somatic mutation and myeloproliferative neoplasm: A Mendelian randomization study. <i>EClinicalMedicine</i> , 2020 , 21, 100280	11.3	13
231	Breast cancer risk factors and their effects on survival: a Mendelian randomisation study. <i>BMC Medicine</i> , 2020 , 18, 327	11.4	7
230	Burden of prediabetes, undiagnosed, and poorly or potentially sub-controlled diabetes: Lolland-Falster health study. <i>BMC Public Health</i> , 2020 , 20, 1711	4.1	2
229	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020 , 107, 837-848	11	12
228	Observational and genetic studies of short telomeres and Alzheimer's disease in 67,000 and 152,000 individuals: a Mendelian randomization study. <i>European Journal of Epidemiology</i> , 2020 , 35, 147-156 ^{12,1}	12.1	12
227	Clinical value of serum hyaluronan and propeptide of type III collagen in patients with pancreatic cancer. <i>International Journal of Cancer</i> , 2020 , 146, 2913-2922	7.5	19
226	Association Analysis of Driver Gene-Related Genetic Variants Identified Novel Lung Cancer Susceptibility Loci with 20,871 Lung Cancer Cases and 15,971 Controls. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020 , 29, 1423-1429	4	2
225	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , 2019 , 9, 12524	4.9	2
224	Appraising the causal relevance of DNA methylation for risk of lung cancer. <i>International Journal of Epidemiology</i> , 2019 , 48, 1493-1504	7.8	27
223	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019 , 10, 431	17.4	45
222	Germline BRCA2 K3326X and CHEK2 I157T mutations increase risk for sporadic pancreatic ductal adenocarcinoma. <i>International Journal of Cancer</i> , 2019 , 145, 686-693	7.5	15

221	Elevated Platelet Count Appears to Be Causally Associated with Increased Risk of Lung Cancer: A Mendelian Randomization Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019 , 28, 935-942	4	12
220	Physical activity and risk of instant and 28-day case-fatality in myocardial infarction. <i>PLoS ONE</i> , 2019 , 14, e0217398	3.7	3
219	Smoking does not accelerate leucocyte telomere attrition: a meta-analysis of 18 longitudinal cohorts. <i>Royal Society Open Science</i> , 2019 , 6, 190420	3.3	16
218	Genetic interaction analysis among oncogenesis-related genes revealed novel genes and networks in lung cancer development. <i>Oncotarget</i> , 2019 , 10, 1760-1774	3.3	12
217	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019 , 10, 1741	17.4	47
216	Smoking and Increased White and Red Blood Cells. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2019 , 39, 965-977	9.4	46
215	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019 , 120, 647-657	8.7	28
214	AHRR (cg05575921) methylation extent of leukocyte DNA and lung cancer survival. <i>PLoS ONE</i> , 2019 , 14, e0211745	3.7	5
213	Arterial and venous thrombosis by high platelet count and high hematocrit: 108521 individuals from the Copenhagen General Population Study. <i>Journal of Thrombosis and Haemostasis</i> , 2019 , 17, 1898-1911	15.1	23
212	Lung Cancer Risk in Never-Smokers of European Descent is Associated With Genetic Variation in the 515.33 TERT-CLPTM1LL Region. <i>Journal of Thoracic Oncology</i> , 2019 , 14, 1360-1369	8.9	12
211	Investigation of Leukocyte Telomere Length and Genetic Variants in Chromosome 5p15.33 as Prognostic Markers in Lung Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019 , 28, 1228-1237	4	5
210	Measured and genetically predicted plasma YKL-40 levels and melanoma mortality. <i>European Journal of Cancer</i> , 2019 , 121, 74-84	7.5	1
209	The p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , 2019 , 5, 38	7.8	12
208	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	3.2	23
207	Prediction and clinical utility of a contralateral breast cancer risk model. <i>Breast Cancer Research</i> , 2019 , 21, 144	8.3	11
206	JAK2-tree: a simple CBC-based decision rule to guide appropriate JAK2 V617F mutation testing. <i>Journal of Clinical Pathology</i> , 2019 , 72, 172-176	3.9	4
205	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019 , 104, 21-34	11	363
204	Bone marrow mononuclear cell telomere length in acute myeloid leukaemia and high-risk myelodysplastic syndrome. <i>European Journal of Haematology</i> , 2019 , 102, 218-226	3.8	4

203	Systematic analyses of regulatory variants in DNase I hypersensitive sites identified two novel lung cancer susceptibility loci. <i>Carcinogenesis</i> , 2019 , 40, 432-440	4.6	3
202	Mononuclear Cell Telomere Attrition Is Associated with Overall Survival after Nonmyeloablative Allogeneic Hematopoietic Cell Transplantation for Hematologic Malignancies. <i>Biology of Blood and Marrow Transplantation</i> , 2019 , 25, 496-504	4.7	2
201	Mendelian Randomization and mediation analysis of leukocyte telomere length and risk of lung and head and neck cancers. <i>International Journal of Epidemiology</i> , 2019 , 48, 751-766	7.8	14
200	Is high vitamin B12 status a cause of lung cancer?. <i>International Journal of Cancer</i> , 2019 , 145, 1499-1503	7.5	33
199	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	7.8	52
198	The BRCA2 c.68-7T>A variant is not pathogenic: A model for clinical calibration of spliceogenicity. <i>Human Mutation</i> , 2018 , 39, 729-741	4.7	16
197	Is smoking heaviness causally associated with alcohol use? A Mendelian randomization study in four European cohorts. <i>International Journal of Epidemiology</i> , 2018 , 47, 1098-1105	7.8	13
196	Genome-wide interaction study of smoking behavior and non-small cell lung cancer risk in Caucasian population. <i>Carcinogenesis</i> , 2018 , 39, 336-346	4.6	11
195	hypomethylation, lung function, lung function decline and respiratory symptoms. <i>European Respiratory Journal</i> , 2018 , 51,	13.6	22
194	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	7.8	53
193	Identification of susceptibility pathways for the role of chromosome 15q25.1 in modifying lung cancer risk. <i>Nature Communications</i> , 2018 , 9, 3221	17.4	29
192	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	36.3	101
191	Serum Biomarker Signature-Based Liquid Biopsy for Diagnosis of Early-Stage Pancreatic Cancer. <i>Journal of Clinical Oncology</i> , 2018 , 36, 2887-2894	2.2	76
190	Fine mapping of MHC region in lung cancer highlights independent susceptibility loci by ethnicity. <i>Nature Communications</i> , 2018 , 9, 3927	17.4	24
189	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	11.6	74
188	Genetic modifiers of radon-induced lung cancer risk: a genome-wide interaction study in former uranium miners. <i>International Archives of Occupational and Environmental Health</i> , 2018 , 91, 937-950	3.2	17
187	Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017 , 77, 2789-2799	10.1	49
186	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	36.3	257

185	TP53 Arg72Pro, mortality after cancer, and all-cause mortality in 105,200 individuals. <i>Scientific Reports</i> , 2017 , 7, 336	4.9	7
184	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	6.6	43
183	Asthma, other atopic conditions and risk of infections in 105 519 general population never and ever smokers. <i>Journal of Internal Medicine</i> , 2017 , 282, 254-267	10.8	14
182	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	36.3	246
181	(cg05575921) hypomethylation marks smoking behaviour, morbidity and mortality. <i>Thorax</i> , 2017 , 72, 646-653	7.3	82
180	Telomere length and depression: prospective cohort study and Mendelian randomisation study in 67 306 individuals. <i>British Journal of Psychiatry</i> , 2017 , 210, 31-38	5.4	19
179	Kringle IV Type 2, Not Low Lipoprotein(a), as a Cause of Diabetes: A Novel Genetic Approach Using SNPs Associated Selectively with Lipoprotein(a) Concentrations or with Kringle IV Type 2 Repeats. <i>Clinical Chemistry</i> , 2017 , 63, 1866-1876	5.5	16
178	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017 , 551, 92-94	50.4	643
177	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017 , 49, 1767-1778	36.3	186
176	Pleiotropy of genetic variants on obesity and smoking phenotypes: Results from the Oncoarray Project of The International Lung Cancer Consortium. <i>PLoS ONE</i> , 2017 , 12, e0185660	3.7	7
175	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. <i>Breast Cancer Research</i> , 2017 , 19, 119	8.3	26
174	Gene-environment interactions involving functional variants: Results from the Breast Cancer Association Consortium. <i>International Journal of Cancer</i> , 2017 , 141, 1830-1840	7.5	13
173	Common breast cancer risk alleles and risk assessment: a study on 35441 individuals from the Danish general population. <i>Annals of Oncology</i> , 2017 , 28, 175-181	10.3	6
172	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017 , 19, 599-603	6.3	51
171	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017 , 26, 126-135	4	183
170	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017 , 46, 1814-1822	7.8	27
169	The prognostic value of serum CA 19-9 in patients with metastatic colorectal cancer.. <i>Journal of Clinical Oncology</i> , 2017 , 35, e15131-e15131	2.2	1
168	Obesity, metabolic factors and risk of different histological types of lung cancer: A Mendelian randomization study. <i>PLoS ONE</i> , 2017 , 12, e0177875	3.7	56

167	- a novel candidate breast cancer susceptibility locus on 6q14.1. <i>Oncotarget</i> , 2017 , 8, 102769-102782	3.3	3
166	The prognostic value of serum IL-6 and YKL-40 in patients with metastatic colorectal cancer.. <i>Journal of Clinical Oncology</i> , 2017 , 35, e15060-e15060	2.2	
165	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016 , 141, 386-401	4.9	15
164	An intergenic risk locus containing an enhancer deletion in 2q35 modulates breast cancer risk by deregulating IGFBP5 expression. <i>Human Molecular Genetics</i> , 2016 , 25, 3863-3876	5.6	24
163	Long telomeres and cancer risk among 95 568 individuals from the general population. <i>International Journal of Epidemiology</i> , 2016 , 45, 1634-1643	7.8	62
162	rs2735383, located at a microRNA binding site in the 3'UTR of NBS1, is not associated with breast cancer risk. <i>Scientific Reports</i> , 2016 , 6, 36874	4.9	2
161	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, 1052-67	24.4	104
160	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016 , 7, 11375	17.4	64
159	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016 , 7, 12675	17.4	53
158	Fine scale mapping of the 17q22 breast cancer locus using dense SNPs, genotyped within the Collaborative Oncological Gene-Environment Study (COGs). <i>Scientific Reports</i> , 2016 , 6, 32512	4.9	16
157	Short Telomere Length and Ischemic Heart Disease: Observational and Genetic Studies in 290 022 Individuals. <i>Clinical Chemistry</i> , 2016 , 62, 1140-9	5.5	70
156	Age- and Tumor Subtype-Specific Breast Cancer Risk Estimates for CHEK2*1100delC Carriers. <i>Journal of Clinical Oncology</i> , 2016 , 34, 2750-60	2.2	107
155	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2016 , 23, 77-91	5.7	41
154	Observationally and Genetically High YKL-40 and Risk of Venous Thromboembolism in the General Population: Cohort and Mendelian Randomization Studies. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2016 , 36, 1030-6	9.4	9
153	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	5.6	55
152	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, 1208-16	2.2	70
151	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	5.8	83
150	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016 , 48, 374-86	36.3	93

149	Genetic variation in the immunosuppression pathway genes and breast cancer susceptibility: a pooled analysis of 42,510 cases and 40,577 controls from the Breast Cancer Association Consortium. <i>Human Genetics</i> , 2016 , 135, 137-54	6.3	6
148	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. <i>Journal of the National Cancer Institute</i> , 2016 , 108,	9.7	65
147	RAD51B in Familial Breast Cancer. <i>PLoS ONE</i> , 2016 , 11, e0153788	3.7	18
146	Association of breast cancer risk with genetic variants showing differential allelic expression: Identification of a novel breast cancer susceptibility locus at 4q21. <i>Oncotarget</i> , 2016 , 7, 80140-80163	3.3	21
145	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	11.6	80
144	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. <i>PLoS ONE</i> , 2016 , 11, e0160316	3.7	11
143	Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. <i>International Journal of Cancer</i> , 2016 , 139, 1303-1317	7.5	26
142	Inflammatory biomarkers and risk of cancer in 84,000 individuals from the general population. <i>International Journal of Cancer</i> , 2016 , 139, 1493-500	7.5	51
141	PALB2, CHEK2 and ATM rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016 , 53, 800-811	5.8	121
140	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	8.3	26
139	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. <i>Breast Cancer Research</i> , 2016 , 18, 64	8.3	25
138	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	9.4	28
137	Genetic predisposition to ductal carcinoma in situ of the breast. <i>Breast Cancer Research</i> , 2016 , 18, 22	8.3	31
136	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. <i>Cancer Causes and Control</i> , 2016 , 27, 679-93	2.8	15
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2	Fine-mapping of 150 breast cancer risk regions identifies 178 high confidence target genes		2
1	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses		2