Stig Bojesen

List of Publications by Citations

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	Genome-wide association study identifies novel breast cancer susceptibility loci. *Nature*, **2007**, 447, 1087-93	50.4	1957
291	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. *Nature Genetics*, **2013**, 45, 353-61, 361e1-2	36.3	813
290	Statin use and reduced cancer-related mortality. *New England Journal of Medicine*, **2012**, 367, 1792-802	59.2	669
289	Association analysis identifies 65 new breast cancer risk loci. *Nature*, **2017**, 551, 92-94	50.4	643
288	A common coding variant in CASP8 is associated with breast cancer risk. *Nature Genetics*, **2007**, 39, 352-8	36.3	557
287	Associations of breast cancer risk factors with tumor subtypes: a pooled analysis from the Breast Cancer Association Consortium studies. *Journal of the National Cancer Institute*, **2011**, 103, 250-63	9.7	513
286	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. *Nature Genetics*, **2013**, 45, 371-84, 384e1-2	36.3	422
285	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. *Nature Genetics*, **2013**, 45, 385-91, 391e1-2	36.3	413
284	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. *Nature Genetics*, **2015**, 47, 373-80	36.3	406
283	Parent-of-origin-specific allelic associations among 106 genomic loci for age at menarche. *Nature*, **2014**, 514, 92-97	50.4	401
282	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. *Nature Genetics*, **2009**, 41, 585-90	36.3	393
281	C-reactive protein as a predictor of prognosis in chronic obstructive pulmonary disease. *American Journal of Respiratory and Critical Care Medicine*, **2007**, 175, 250-5	10.2	391
280	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. *American Journal of Human Genetics*, **2019**, 104, 21-34	11	363
279	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. *Nature Genetics*, **2013**, 45, 392-8, 398e1-2	36.3	327
278	Prediction of breast cancer risk based on profiling with common genetic variants. *Journal of the National Cancer Institute*, **2015**, 107,	9.7	324
277	MicroRNA biomarkers in whole blood for detection of pancreatic cancer. *JAMA - Journal of the American Medical Association*, **2014**, 311, 392-404	27.4	315
276	Baseline C-reactive protein is associated with incident cancer and survival in patients with cancer. *Journal of Clinical Oncology*, **2009**, 27, 2217-24	2.2	305

275	Heterogeneity of breast cancer associations with five susceptibility loci by clinical and pathological characteristics. *PLoS Genetics*, **2008**, 4, e1000054	6	280
274	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. *Nature Genetics*, **2017**, 49, 834-841	36.3	257
273	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. *Nature Genetics*, **2017**, 49, 1126-1132	36.3	246
272	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. *Nature Genetics*, **2011**, 43, 785-91	36.3	243
271	Genome-wide association analysis identifies three new breast cancer susceptibility loci. *Nature Genetics*, **2012**, 44, 312-8	36.3	237
270	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. *Nature Genetics*, **2015**, 47, 1294-1303	36.3	226
269	CHEK2*1100delC genotyping for clinical assessment of breast cancer risk: meta-analyses of 26,000 patient cases and 27,000 controls. *Journal of Clinical Oncology*, **2008**, 26, 542-8	2.2	216
268	Peripheral blood leukocyte telomere length and mortality among 64,637 individuals from the general population. *Journal of the National Cancer Institute*, **2015**, 107, djv074	9.7	200
267	Low 25-hydroxyvitamin D and risk of type 2 diabetes: a prospective cohort study and metaanalysis. *Clinical Chemistry*, **2013**, 59, 381-91	5.5	189
266	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. *Nature Genetics*, **2017**, 49, 1767-1778	36.3	186
265	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. *Cancer Epidemiology Biomarkers and Prevention*, **2017**, 26, 126-135	4	183
264	Genetically low vitamin D concentrations and increased mortality: Mendelian randomisation analysis in three large cohorts. *BMJ, The*, **2014**, 349, g6330	5.9	182
263	Short telomere length, cancer survival, and cancer risk in 47102 individuals. *Journal of the National Cancer Institute*, **2013**, 105, 459-68	9.7	168
262	Functional variants at the 11q13 risk locus for breast cancer regulate cyclin D1 expression through long-range enhancers. *American Journal of Human Genetics*, **2013**, 92, 489-503	11	167
261	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. *Lancet, The*, **2021**, 398, 957-980	40	154
260	Short telomere length, myocardial infarction, ischemic heart disease, and early death. *Arteriosclerosis, Thrombosis, and Vascular Biology*, **2012**, 32, 822-9	9.4	152
259	Breast Cancer Risk Genes - Association Analysis in More than 113,000 Women. *New England Journal of Medicine*, **2021**, 384, 428-439	59.2	143
258	Low penetrance breast cancer susceptibility loci are associated with specific breast tumor subtypes: findings from the Breast Cancer Association Consortium. *Human Molecular Genetics*, **2011** , 20, 3289-303	5.6	140

257	CHEK2*1100delC heterozygosity in women with breast cancer associated with early death, breast cancer-specific death, and increased risk of a second breast cancer. *Journal of Clinical Oncology*, **2012**, 30, 4308-16	2.2	134
256	PALB2, CHEK2 and ATM rare variants and cancer risk: data from COGS. *Journal of Medical Genetics*, **2016**, 53, 800-811	5.8	121
255	Vitamin D concentration, obesity, and risk of diabetes: a mendelian randomisation study. *Lancet Diabetes and Endocrinology,the*, **2014**, 2, 298-306	18.1	120
254	Short telomere length, lung function and chronic obstructive pulmonary disease in 46,396 individuals. *Thorax*, **2013**, 68, 429-35	7.3	118
253	Telomere shortening unrelated to smoking, body weight, physical activity, and alcohol intake: 4,576 general population individuals with repeat measurements 10 years apart. *PLoS Genetics*, **2014**, 10, e1004191	6	112
252	Evidence of gene-environment interactions between common breast cancer susceptibility loci and established environmental risk factors. *PLoS Genetics*, **2013**, 9, e1003284	6	112
251	Reduced 25-hydroxyvitamin D and risk of Alzheimer's disease and vascular dementia. *Alzheimers and Dementia*, **2014**, 10, 296-302	1.2	111
250	A meta-analysis of genome-wide association studies to identify prostate cancer susceptibility loci associated with aggressive and non-aggressive disease. *Human Molecular Genetics*, **2013**, 22, 408-15	5.6	109
249	Age- and Tumor Subtype-Specific Breast Cancer Risk Estimates for CHEK2*1100delC Carriers. *Journal of Clinical Oncology*, **2016**, 34, 2750-60	2.2	107
248	A genome-wide association scan (GWAS) for mean telomere length within the COGS project: identified loci show little association with hormone-related cancer risk. *Human Molecular Genetics*, **2013**, 22, 5056-64	5.6	107
247	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. *Cancer Discovery*, **2016**, 6, 1052-67	24.4	104
246	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. *Nature Genetics*, **2018**, 50, 968-978	36.3	101
245	Tumor suppressor p53 Arg72Pro polymorphism and longevity, cancer survival, and risk of cancer in the general population. *Journal of Experimental Medicine*, **2007**, 204, 1295-301	16.6	100
244	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. *Nature Genetics*, **2016**, 48, 374-86	36.3	93
243	19p13.1 is a triple-negative-specific breast cancer susceptibility locus. *Cancer Research*, **2012**, 72, 1795-803	10.1	93
242	Identification of a BRCA2-specific modifier locus at 6p24 related to breast cancer risk. *PLoS Genetics*, **2013**, 9, e1003173	6	90
241	Plasma YKL-40 levels in healthy subjects from the general population. *Clinica Chimica Acta*, **2011**, 412, 709-12	6.2	90
240	C-reactive protein and the risk of cancer: a mendelian randomization study. *Journal of the National Cancer Institute*, **2010**, 102, 202-6	9.7	90

239	Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with TERT expression. *Human Molecular Genetics*, **2013**, 22, 2520-8	5.6	88
238	Increased risk of breast cancer associated with CHEK2*1100delC. *Journal of Clinical Oncology*, **2007**, 25, 57-63	2.2	88
237	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. *Nature Communications*, **2014**, 4, 4999	17.4	87
236	The JAK2 V617F somatic mutation, mortality and cancer risk in the general population. *Haematologica*, **2011**, 96, 450-3	6.6	85
235	Telomeres and human health. *Journal of Internal Medicine*, **2013**, 274, 399-413	10.8	84
234	Genome-wide association study identifies a common variant in RAD51B associated with male breast cancer risk. *Nature Genetics*, **2012**, 44, 1182-4	36.3	84
233	No evidence that protein truncating variants in BRIP1 are associated with breast cancer risk: implications for gene panel testing. *Journal of Medical Genetics*, **2016**, 53, 298-309	5.8	83
232	(cg05575921) hypomethylation marks smoking behaviour, morbidity and mortality. *Thorax*, **2017**, 72, 646-653	7.3	82
231	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	8.3	82
230	Fine-scale mapping of the FGFR2 breast cancer risk locus: putative functional variants differentially bind FOXA1 and E2F1. *American Journal of Human Genetics*, **2013**, 93, 1046-60	11	80
229	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. *PLoS Medicine*, **2016**, 13, e1002105	11.6	80
228	Risk of cancer among HIV-infected individuals compared to the background population: impact of smoking and HIV. *Aids*, **2014**, 28, 1499-508	3.5	77
227	Low plasma 25-hydroxyvitamin D and risk of tobacco-related cancer. *Clinical Chemistry*, **2013**, 59, 771-80	5.5	77
226	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. *Nature Genetics*, **2020**, 52, 572-581	36.3	76
225	Serum Biomarker Signature-Based Liquid Biopsy for Diagnosis of Early-Stage Pancreatic Cancer. *Journal of Clinical Oncology*, **2018**, 36, 2887-2894	2.2	76
224	Association of ESR1 gene tagging SNPs with breast cancer risk. *Human Molecular Genetics*, **2009**, 18, 1131-9	5.6	75
223	The role of genetic breast cancer susceptibility variants as prognostic factors. *Human Molecular Genetics*, **2012**, 21, 3926-39	5.6	75
222	Height and Breast Cancer Risk: Evidence From Prospective Studies and Mendelian Randomization. *Journal of the National Cancer Institute*, **2015**, 107,	9.7	74

221	Lymphopenia and risk of infection and infection-related death in 98,344 individuals from a prospective Danish population-based study. *PLoS Medicine*, **2018**, 15, e1002685	11.6	74
220	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. *Lancet, The*, **2020**, 396, 1511-1524	40	73
219	Short Telomere Length and Ischemic Heart Disease: Observational and Genetic Studies in 290 022 Individuals. *Clinical Chemistry*, **2016**, 62, 1140-9	5.5	70
218	Increased Risk for Other Cancers in Addition to Breast Cancer for CHEK2*1100delC Heterozygotes Estimated From the Copenhagen General Population Study. *Journal of Clinical Oncology*, **2016**, 34, 1208-16	2.2	70
217	Elevated plasma YKL-40 predicts increased risk of gastrointestinal cancer and decreased survival after any cancer diagnosis in the general population. *Journal of Clinical Oncology*, **2009**, 27, 572-8	2.2	70
216	Diagnostic value of JAK2 V617F somatic mutation for myeloproliferative cancer in 49 488 individuals from the general population. *British Journal of Haematology*, **2013**, 160, 70-9	4.5	68
215	The common germline Arg72Pro polymorphism of p53 and increased longevity in humans. *Cell Cycle*, **2008**, 7, 158-63	4.7	68
214	Prevalence of the HOXB13 G84E germline mutation in British men and correlation with prostate cancer risk, tumour characteristics and clinical outcomes. *Annals of Oncology*, **2015**, 26, 756-761	10.3	67
213	Associations of common variants at 1p11.2 and 14q24.1 (RAD51L1) with breast cancer risk and heterogeneity by tumor subtype: findings from the Breast Cancer Association Consortium. *Human Molecular Genetics*, **2011**, 20, 4693-706	5.6	66
212	BRCA2 Polymorphic Stop Codon K3326X and the Risk of Breast, Prostate, and Ovarian Cancers. *Journal of the National Cancer Institute*, **2016**, 108,	9.7	65
211	Increased body mass index, elevated C-reactive protein, and short telomere length. *Journal of Clinical Endocrinology and Metabolism*, **2014**, 99, E1671-5	5.6	65
210	Integrin beta3 Leu33Pro homozygosity and risk of cancer. *Journal of the National Cancer Institute*, **2003**, 95, 1150-7	9.7	65
209	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. *Nature Communications*, **2016**, 7, 11375	17.4	64
208	Plasma 25-hydroxyvitamin D, lung function and risk of chronic obstructive pulmonary disease. *Thorax*, **2014**, 69, 24-31	7.3	63
207	Long telomeres and cancer risk among 95 568 individuals from the general population. *International Journal of Epidemiology*, **2016**, 45, 1634-1643	7.8	62
206	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. *Nature Genetics*, **2021**, 53, 65-75	36.3	62
205	Association of clinical benign prostate hyperplasia with prostate cancer incidence and mortality revisited: a nationwide cohort study of 3,009,258 men. *European Urology*, **2011**, 60, 691-8	10.2	61
204	Fine-scale mapping of the 5q11.2 breast cancer locus reveals at least three independent risk variants regulating MAP3K1. *American Journal of Human Genetics*, **2015**, 96, 5-20	11	59

203	Statin use and reduced cancer-related mortality. *New England Journal of Medicine*, **2013**, 368, 576-7	59.2	58
202	Plasma YKL-40 and total and disease-specific mortality in the general population. *Clinical Chemistry*, **2010**, 56, 1580-91	5.5	56
201	Elevated plasma YKL-40 levels and ischemic stroke in the general population. *Annals of Neurology*, **2010**, 68, 672-80	9.4	56
200	Obesity, metabolic factors and risk of different histological types of lung cancer: A Mendelian randomization study. *PLoS ONE*, **2017**, 12, e0177875	3.7	56
199	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. *Nature Genetics*, **2020**, 52, 56-73	36.3	56
198	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. *Human Molecular Genetics*, **2016**, 25, 2256-2268	5.6	55
197	Platelet glycoprotein IIb/IIIa Pl(A2)/Pl(A2) homozygosity associated with risk of ischemic cardiovascular disease and myocardial infarction in young men: the Copenhagen City Heart Study. *Journal of the American College of Cardiology*, **2003**, 42, 661-7	15.1	55
196	Multiple novel prostate cancer susceptibility signals identified by fine-mapping of known risk loci among Europeans. *Human Molecular Genetics*, **2015**, 24, 5589-602	5.6	54
195	JAK2V617F somatic mutation in the general population: myeloproliferative neoplasm development and progression rate. *Haematologica*, **2014**, 99, 1448-55	6.6	54
194	Joint associations of a polygenic risk score and environmental risk factors for breast cancer in the Breast Cancer Association Consortium. *International Journal of Epidemiology*, **2018**, 47, 526-536	7.8	53
193	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. *Nature Communications*, **2016**, 7, 12675	17.4	53
192	Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis. *International Journal of Epidemiology*, **2019**, 48, 795-806	7.8	52
191	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. *Genetics in Medicine*, **2017**, 19, 599-603	6.8	51
190	Inflammatory biomarkers and risk of cancer in 84,000 individuals from the general population. *International Journal of Cancer*, **2016**, 139, 1493-500	7.5	51
189	Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. *Cancer Research*, **2017**, 77, 2789-2799	10.1	49
188	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. *Human Molecular Genetics*, **2014**, 23, 6096-111	5.6	48
187	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. *Nature Communications*, **2019**, 10, 1741	17.4	47
186	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. *International Journal of Cancer*, **2011**, 128, 644-52	7.5	47

185	Smoking and Increased White and Red Blood Cells. *Arteriosclerosis, Thrombosis, and Vascular Biology* , **2019**, 39, 965-977	9.4	46
184	Shared heritability and functional enrichment across six solid cancers. *Nature Communications*, **2019** , 10, 431	17.4	45
183	Association between a germline OCA2 polymorphism at chromosome 15q13.1 and estrogen receptor-negative breast cancer survival. *Journal of the National Cancer Institute*, **2010**, 102, 650-62	9.7	45
182	Nicotinic acetylcholine receptor polymorphism, smoking behavior, and tobacco-related cancer and lung and cardiovascular diseases: a cohort study. *Journal of Clinical Oncology*, **2011**, 29, 2875-82	2.2	44
181	Shorter leukocyte telomere length is associated with higher risk of infections: a prospective study of 75,309 individuals from the general population. *Haematologica*, **2017**, 102, 1457-1465	6.6	43
180	Evidence that the 5p12 Variant rs10941679 Confers Susceptibility to Estrogen-Receptor-Positive Breast Cancer through FGF10 and MRPS30 Regulation. *American Journal of Human Genetics*, **2016**, 99, 903-911	11	43
179	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. *Endocrine-Related Cancer*, **2016**, 23, 77-91	5.7	41
178	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	4.3	40
177	Identification of novel genetic markers of breast cancer survival. *Journal of the National Cancer Institute*, **2015**, 107,	9.7	38
176	Genetic predisposition to in situ and invasive lobular carcinoma of the breast. *PLoS Genetics*, **2014**, 10, e1004285	6	38
175	MicroRNA related polymorphisms and breast cancer risk. *PLoS ONE*, **2014**, 9, e109973	3.7	37
174	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. *Human Molecular Genetics*, **2015**, 24, 2966-84	5.6	36
173	CHRNA3 genotype, nicotine dependence, lung function and disease in the general population. *European Respiratory Journal*, **2012**, 40, 1538-44	13.6	36
172	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. *Human Molecular Genetics*, **2015**, 24, 285-98	5.6	35
171	DNA mismatch repair gene MSH6 implicated in determining age at natural menopause. *Human Molecular Genetics*, **2014**, 23, 2490-7	5.6	35
170	11q13 is a susceptibility locus for hormone receptor positive breast cancer. *Human Mutation*, **2012**, 33, 1123-32	4.7	33
169	Is high vitamin B12 status a cause of lung cancer?. *International Journal of Cancer*, **2019**, 145, 1499-1503	7.5	33
168	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, 1473-83	7.8	32

167	Prostate-specific antigen and long-term prediction of prostate cancer incidence and mortality in the general population. *European Urology*, **2012**, 61, 865-74	10.2	32
166	Copy number variation in glutathione-S-transferase T1 and M1 predicts incidence and 5-year survival from prostate and bladder cancer, and incidence of corpus uteri cancer in the general population. *Pharmacogenomics Journal*, **2011**, 11, 292-9	3.5	32
165	Missense variants in ATM in 26,101 breast cancer cases and 29,842 controls. *Cancer Epidemiology Biomarkers and Prevention*, **2010**, 19, 2143-51	4	31
164	Genetic predisposition to ductal carcinoma in situ of the breast. *Breast Cancer Research*, **2016**, 18, 22	8.3	31
163	Elevated plasma YKL-40, lipids and lipoproteins, and ischemic vascular disease in the general population. *Stroke*, **2015**, 46, 329-35	6.7	29
162	Identification of susceptibility pathways for the role of chromosome 15q25.1 in modifying lung cancer risk. *Nature Communications*, **2018**, 9, 3221	17.4	29
161	Alcohol consumption and survival after a breast cancer diagnosis: a literature-based meta-analysis and collaborative analysis of data for 29,239 cases. *Cancer Epidemiology Biomarkers and Prevention*, **2014**, 23, 934-45	4	29
160	Genome-wide association study of germline variants and breast cancer-specific mortality. *British Journal of Cancer*, **2019**, 120, 647-657	8.7	28
159	A large-scale assessment of two-way SNP interactions in breast cancer susceptibility using 46,450 cases and 42,461 controls from the breast cancer association consortium. *Human Molecular Genetics*, **2014**, 23, 1934-46	5.6	28
158	Plasma testosterone in the general population, cancer prognosis and cancer risk: a prospective cohort study. *Annals of Oncology*, **2014**, 25, 712-718	10.3	28
157	Role of inflammatory marker YKL-40 in the diagnosis, prognosis and cause of cardiovascular and liver diseases. *Critical Reviews in Clinical Laboratory Sciences*, **2016**, 53, 396-408	9.4	28
156	Genetic insights into biological mechanisms governing human ovarian ageing. *Nature*, **2021**, 596, 393-397	50.4	28
155	Appraising the causal relevance of DNA methylation for risk of lung cancer. *International Journal of Epidemiology*, **2019**, 48, 1493-1504	7.8	27
154	Body mass index and breast cancer survival: a Mendelian randomization analysis. *International Journal of Epidemiology*, **2017**, 46, 1814-1822	7.8	27
153	Confirmation of 5p12 as a susceptibility locus for progesterone-receptor-positive, lower grade breast cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2011**, 20, 2222-31	4	27
152	Missense polymorphisms in BRCA1 and BRCA2 and risk of breast and ovarian cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2009**, 18, 2339-42	4	27
151	Cancer risk by combined levels of YKL-40 and C-reactive protein in the general population. *British Journal of Cancer*, **2012**, 106, 199-205	8.7	27
150	Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression. *American Journal of Human Genetics*, **2015**, 97, 22-34	11	26

149	Multi-stage genome-wide association study identifies new susceptibility locus for testicular germ cell tumour on chromosome 3q25. *Human Molecular Genetics*, **2015**, 24, 1169-76	5.6	26
148	Reproductive profiles and risk of breast cancer subtypes: a multi-center case-only study. *Breast Cancer Research*, **2017**, 19, 119	8.3	26
147	Investigation of gene-environment interactions between 47 newly identified breast cancer susceptibility loci and environmental risk factors. *International Journal of Cancer*, **2015**, 136, E685-96	7.5	26
146	IgE and risk of cancer in 37 747 individuals from the general population. *Annals of Oncology*, **2015**, 26, 1784-90	10.3	26
145	Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. *International Journal of Cancer*, **2016**, 139, 1303-1317	7.5	26
144	Patient survival and tumor characteristics associated with CHEK2:p.I157T - findings from the Breast Cancer Association Consortium. *Breast Cancer Research*, **2016**, 18, 98	8.3	26
143	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. *Breast Cancer Research*, **2016**, 18, 64	8.3	25
142	An intergenic risk locus containing an enhancer deletion in 2q35 modulates breast cancer risk by deregulating IGFBP5 expression. *Human Molecular Genetics*, **2016**, 25, 3863-3876	5.6	24
141	Identification of new genetic susceptibility loci for breast cancer through consideration of gene-environment interactions. *Genetic Epidemiology*, **2014**, 38, 84-93	2.6	24
140	Common germline polymorphisms associated with breast cancer-specific survival. *Breast Cancer Research*, **2015**, 17, 58	8.3	24
139	Fine mapping of MHC region in lung cancer highlights independent susceptibility loci by ethnicity. *Nature Communications*, **2018**, 9, 3927	17.4	24
138	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	15.4	23
137	Diagnostic and Prognostic Impact of Circulating YKL-40, IL-6, and CA 19.9 in Patients with Pancreatic Cancer. *PLoS ONE*, **2013**, 8, e67059	3.7	23
136	Secular trends in smoking in relation to prevalent and incident smoking-related disease: A prospective population-based study. *Tobacco Induced Diseases*, **2019**, 17, 72	3.2	23
135	The potential diagnostic value of serum microRNA signature in patients with pancreatic cancer. *International Journal of Cancer*, **2016**, 139, 2312-24	7.5	23
134	hypomethylation, lung function, lung function decline and respiratory symptoms. *European Respiratory Journal*, **2018**, 51,	13.6	22
133	Cost-effectiveness of surveillance programs for families at high and moderate risk of hereditary non-polyposis colorectal cancer. *International Journal of Technology Assessment in Health Care*, **2007**, 23, 89-95	1.8	22
132	Association of breast cancer risk with genetic variants showing differential allelic expression: Identification of a novel breast cancer susceptibility locus at 4q21. *Oncotarget*, **2016**, 7, 80140-80163	3.3	21

131	Detection and characterization of lung cancer using cell-free DNA fragmentomes. *Nature Communications*, **2021**, 12, 5060	17.4	21
130	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. *Nature Communications*, **2020**, 11, 312	17.4	20
129	FGF receptor genes and breast cancer susceptibility: results from the Breast Cancer Association Consortium. *British Journal of Cancer*, **2014**, 110, 1088-100	8.7	20
128	Characterisation of the GRAF gene promoter and its methylation in patients with acute myeloid leukaemia and myelodysplastic syndrome. *British Journal of Cancer*, **2006**, 94, 323-32	8.7	20
127	Smoking, blood cells and myeloproliferative neoplasms: meta-analysis and Mendelian randomization of 2 million people. *British Journal of Haematology*, **2020**, 189, 323-334	4.5	20
126	Telomere length and depression: prospective cohort study and Mendelian randomisation study in 67 306 individuals. *British Journal of Psychiatry*, **2017**, 210, 31-38	5.4	19
125	SNP-SNP interaction analysis of NF- signaling pathway on breast cancer survival. *Oncotarget*, **2015**, 6, 37979-94	3.3	19
124	Common polymorphisms in CYP2C9, subclinical atherosclerosis and risk of ischemic vascular disease in 52,000 individuals. *Pharmacogenomics Journal*, **2009**, 9, 327-32	3.5	19
123	Clinical value of serum hyaluronan and propeptide of type III collagen in patients with pancreatic cancer. *International Journal of Cancer*, **2020**, 146, 2913-2922	7.5	19
122	RAD51B in Familial Breast Cancer. *PLoS ONE*, **2016**, 11, e0153788	3.7	18
121	Causal relationships between body mass index, smoking and lung cancer: Univariable and multivariable Mendelian randomization. *International Journal of Cancer*, **2021**, 148, 1077-1086	7.5	18
120	Fine-scale mapping of the 4q24 locus identifies two independent loci associated with breast cancer risk. *Cancer Epidemiology Biomarkers and Prevention*, **2015**, 24, 1680-91	4	17
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