Carl Blomqvist

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68 18,586 287 130 h-index g-index citations papers 5.36 21,736 7.8 300 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
287	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013 , 45, 353-61, 361e1-2	36.3	813
286	53BP1 loss rescues BRCA1 deficiency and is associated with triple-negative and BRCA-mutated breast cancers. <i>Nature Structural and Molecular Biology</i> , 2010 , 17, 688-95	17.6	707
285	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017 , 551, 92-94	50.4	643
284	Subtyping of breast cancer by immunohistochemistry to investigate a relationship between subtype and short and long term survival: a collaborative analysis of data for 10,159 cases from 12 studies. <i>PLoS Medicine</i> , 2010 , 7, e1000279	11.6	616
283	A common coding variant in CASP8 is associated with breast cancer risk. <i>Nature Genetics</i> , 2007 , 39, 352-	836.3	557
282	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, 250-63	9.7	513
281	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013 , 45, 371-84, 384e1-2	36.3	422
280	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015 , 47, 373-80	36.3	406
279	Parent-of-origin-specific allelic associations among 106 genomic loci for age at menarche. <i>Nature</i> , 2014 , 514, 92-97	50.4	401
278	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. <i>Nature Genetics</i> , 2009 , 41, 585-	99 6.3	393
277	Tailored fluorouracil, epirubicin, and cyclophosphamide compared with marrow-supported high-dose chemotherapy as adjuvant treatment for high-risk breast cancer: a randomised trial. Scandinavian Breast Group 9401 study. <i>Lancet, The</i> , 2000 , 356, 1384-91	40	374
276	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
275	A CHEK2 genetic variant contributing to a substantial fraction of familial breast cancer. <i>American Journal of Human Genetics</i> , 2002 , 71, 432-8	11	354
274	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013 , 45, 392-8, 398e1-2	36.3	327
273	Prediction of breast cancer risk based on profiling with common genetic variants. <i>Journal of the National Cancer Institute</i> , 2015 , 107,	9.7	324
272	Heterogeneity of breast cancer associations with five susceptibility loci by clinical and pathological characteristics. <i>PLoS Genetics</i> , 2008 , 4, e1000054	6	280
271	Adjuvant clodronate treatment does not reduce the frequency of skeletal metastases in node-positive breast cancer patients: 5-year results of a randomized controlled trial. <i>Journal of Clinical Oncology</i> , 2001 , 19, 10-7	2.2	278

(2020-2011)

A common variant at the TERT-CLPTM1L locus is associated with estrogen receptor-negative breast cancer. <i>Nature Genetics</i> , 2011 , 43, 1210-4	36.3	253
Distribution of coronary artery stenosis after radiation for breast cancer. <i>Journal of Clinical Oncology</i> , 2012 , 30, 380-6	2.2	250
Genome-wide association analysis identifies three new breast cancer susceptibility loci. <i>Nature Genetics</i> , 2012 , 44, 312-8	36.3	237
Prognostic impact of immunohistochemically defined germinal center phenotype in diffuse large B-cell lymphoma patients treated with immunochemotherapy. <i>Blood</i> , 2007 , 109, 4930-5	2.2	235
The combined status of ATM and p53 link tumor development with therapeutic response. <i>Genes and Development</i> , 2009 , 23, 1895-909	12.6	234
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	36.3	226
Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017 , 49, 1767-1778	36.3	186
Cement is recommended in intralesional surgery of giant cell tumors: a Scandinavian Sarcoma Group study of 294 patients followed for a median time of 5 years. <i>Monthly Notices of the Royal Astronomical Society: Letters</i> , 2008 , 79, 86-93	4.3	177
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	11	167
Ten-year follow-up of a randomized controlled trial of adjuvant clodronate treatment in node-positive breast cancer patients. <i>Acta Oncològica</i> , 2004 , 43, 650-6	3.2	165
NAD(P)H:quinone oxidoreductase 1 NQO1*2 genotype (P187S) is a strong prognostic and predictive factor in breast cancer. <i>Nature Genetics</i> , 2008 , 40, 844-53	36.3	163
A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. <i>Human Molecular Genetics</i> , 2012 , 21, 5373-84	5.6	143
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
Genomic subtypes of breast cancer identified by array-comparative genomic hybridization display distinct molecular and clinical characteristics. <i>Breast Cancer Research</i> , 2010 , 12, R42	8.3	141
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, 3289-303	5.6	140
Tamoxifen treatment after adjuvant chemotherapy has opposite effects on bone mineral density in premenopausal patients depending on menstrual status. <i>Journal of Clinical Oncology</i> , 2006 , 24, 675-80	2.2	137
CHEK2*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, 4308-16	2.2	134
Cancer Risks Associated With Germline Pathogenic Variants: An International Study of 524 Families. Journal of Clinical Oncology, 2020 , 38, 674-685	2.2	133
	Cenome-wide association analysis identifies three new breast cancer. Journal of Clinical Oncology, 2012, 30, 380-6 Genome-wide association analysis identifies three new breast cancer susceptibility loci. Nature Genetics, 2012, 44, 312-8 Prognostic impact of immunohistochemically defined germinal center phenotype in diffuse large B-cell lymphoma patients treated with immunochemotherapy. Blood, 2007, 109, 4930-5 The combined status of ATM and p53 link tumor development with therapeutic response. Genes and Development, 2009, 23, 1895-909 Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. Nature Genetics, 2015, 47, 1294-1303 Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778 Cement is recommended in intralesional surgery of giant cell tumors: a Scandinavian Sarcoma Group study of 294 patients followed for a median time of 5 years. Monthly Notices of the Royal Astronomical Society: Letters, 2008, 79, 86-93 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 Ten-year follow-up of a randomized controlled trial of adjuvant clodronate treatment in node-positive breast cancer patients. Acta Onco@ica, 2004, 43, 650-6 NAD(P)H:quinone oxidoreductase 1 NQO1*2 genotype (P1875) is a strong prognostic and predictive factor in breast cancer. Nature Genetics, 2008, 40, 844-53 A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. Human Molecular Genetics, 2012, 21, 5373-84 Breast Cancer Risk Genes - Association Analysis in More than 113,000 Women. New England Journal of Medicine, 2021, 384, 428-439 Genomic subtypes of breast cancer identified by array-comparative genomic hybridization display distinct molecular and clinical characteristics. Breast Cancer R	Distribution of coronary artery stenosis after radiation for breast cancer. Journal of Clinical Oncology, 2012, 30, 380-6 Genome-wide association analysis identifies three new breast cancer susceptibility loci. Nature Genetics, 2012, 44, 312-8 Prognostic impact of immunohistochemically defined germinal center phenotype in diffuse large B-cell lymphoma patients treated with immunochemotherapy. Blood, 2007, 109, 4930-5 The combined status of ATM and p53 link tumor development with therapeutic response. Genes and Development, 2009, 23, 1895-909 Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. Nature Genetics, 2015, 47, 1294-1303 Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778 Cement is recommended in intralesional surgery of giant cell tumors: a Scandinavian Sarcoma Group study of 294 patients followed for a median time of 5 years. Monthly Notices of the Royal Astronomical Society: Letters, 2008, 79, 86-93 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 NAD(P)H:quinone oxidoreductase 1 NQO1+2 genotype (P187S) is a strong prognostic and predictive factor in breast cancer. Nature Genetics, 2008, 40, 844-53 Ameta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. Human Molecular Genetics, 2012, 21, 5373-84 Breast Cancer Risk Genes - Association Analysis in More than 113,000 Women. New England Journal of Medicine, 2021, 384, 428-439 Genomic subtypes of preast cancer identified by array-comparative genomic hybridization display distinct molecular and clinical characteristics. Breast Cancer Research, 2010, 12, R42 Low penetrance breast cancer susceptibility loci are associated with specific breast tumor subtypes: findings from the Breast Ca

252	Treatment-related factors predisposing to chronic pain in patients with breast cancera multivariate approach. <i>Acta Oncològica</i> , 1997 , 36, 625-30	3.2	132
251	Exome sequencing identifies FANCM as a susceptibility gene for triple-negative breast cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014 , 111, 15172-7	11.5	128
250	Breast cancer patients with p53 Pro72 homozygous genotype have a poorer survival. <i>Clinical Cancer Research</i> , 2005 , 11, 5098-103	12.9	126
249	Genome-wide association study identifies 25 known breast cancer susceptibility loci as risk factors for triple-negative breast cancer. <i>Carcinogenesis</i> , 2014 , 35, 1012-9	4.6	121
248	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
247	Aberrations of the MRE11-RAD50-NBS1 DNA damage sensor complex in human breast cancer: MRE11 as a candidate familial cancer-predisposing gene. <i>Molecular Oncology</i> , 2008 , 2, 296-316	7.9	120
246	CHEK2 variant I157T may be associated with increased breast cancer risk. <i>International Journal of Cancer</i> , 2004 , 111, 543-7	7.5	114
245	DNA sequence copy number increase at 8q: a potential new prognostic marker in high-grade osteosarcoma. <i>International Journal of Cancer</i> , 1999 , 84, 114-21	7.5	114
244	RAD51C is a susceptibility gene for ovarian cancer. <i>Human Molecular Genetics</i> , 2011 , 20, 3278-88	5.6	111
243	The breast cancer susceptibility mutation PALB2 1592delT is associated with an aggressive tumor phenotype. <i>Clinical Cancer Research</i> , 2009 , 15, 3214-22	12.9	106
242	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, 1052-6	3 7 4·4	104
241	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
240	Common breast cancer susceptibility loci are associated with triple-negative breast cancer. <i>Cancer Research</i> , 2011 , 71, 6240-9	10.1	100
239	Clinical course of nonvisceral soft tissue leiomyosarcoma in 225 patients from the Scandinavian Sarcoma Group. <i>Cancer</i> , 2007 , 109, 282-91	6.4	98
238	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
237	19p13.1 is a triple-negative-specific breast cancer susceptibility locus. <i>Cancer Research</i> , 2012 , 72, 1795-8	303.1	93
236	Risk of estrogen receptor-positive and -negative breast cancer and single-nucleotide polymorphism 2q35-rs13387042. <i>Journal of the National Cancer Institute</i> , 2009 , 101, 1012-8	9.7	90
235	The predictive value of bcl-2, bax, bcl-xL, bag-1, fas, and fasL for chemotherapy response in advanced breast cancer. <i>Clinical Cancer Research</i> , 2002 , 8, 811-6	12.9	90

234	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014 , 4, 4999	17.4	87
233	Prognostic role of HuR in hereditary breast cancer. <i>Clinical Cancer Research</i> , 2007 , 13, 6959-63	12.9	86
232	Bisphosphonate therapy in metastatic breast cancer. Acta Oncol@ica, 1996, 35 Suppl 5, 81-3	3.2	86
231	A combined analysis of genome-wide association studies in breast cancer. <i>Breast Cancer Research and Treatment</i> , 2011 , 126, 717-27	4.4	85
230	Correlation of CHEK2 protein expression and c.1100delC mutation status with tumor characteristics among unselected breast cancer patients. <i>International Journal of Cancer</i> , 2005 , 113, 575	5 - 86	85
229	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	8.3	82
228	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, 1046-60	11	80
227	A high serum matrix metalloproteinase-2 level is associated with an adverse prognosis in node-positive breast carcinoma. <i>Clinical Cancer Research</i> , 2004 , 10, 1057-63	12.9	78
226	Familial Breast Cancer in Southern Finland. <i>Disease Markers</i> , 1999 , 15, 105-105	3.2	78
225	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
224	The role of genetic breast cancer susceptibility variants as prognostic factors. <i>Human Molecular Genetics</i> , 2012 , 21, 3926-39	5.6	75
223	Histopathological features of breast tumours in BRCA1, BRCA2 and mutation-negative breast cancer families. <i>Breast Cancer Research</i> , 2005 , 7, R93-100	8.3	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	Gains and losses of DNA sequences in liposarcomas evaluated by comparative genomic hybridization. <i>Genes Chromosomes and Cancer</i> , 1996 , 15, 89-94	5	73
220	Do MDM2 SNP309 and TP53 R72P interact in breast cancer susceptibility? A large pooled series from the breast cancer association consortium. <i>Cancer Research</i> , 2007 , 67, 9584-90	10.1	69
219	Increased incidence of stroke in women with breast cancer. European Journal of Cancer, 2005, 41, 423-9	7.5	67
218	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, 4693-706	5.6	66
217	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

216	MiR-34a expression has an effect for lower risk of metastasis and associates with expression patterns predicting clinical outcome in breast cancer. <i>PLoS ONE</i> , 2011 , 6, e26122	3.7	65
215	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016 , 7, 11375	17.4	64
214	Multiple founder effects and geographical clustering of BRCA1 and BRCA2 families in Finland. <i>European Journal of Human Genetics</i> , 2000 , 8, 757-63	5.3	64
213	A genome-wide association study of early-onset breast cancer identifies PFKM as a novel breast cancer gene and supports a common genetic spectrum for breast cancer at any age. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014 , 23, 658-69	4	63
212	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	11	59
211	Overrepresentation of 1q21-23 and 12q13-21 in lipoma-like liposarcomas but not in benign lipomas: a comparative genomic hybridization study. <i>Cancer Genetics and Cytogenetics</i> , 1997 , 99, 14-8		59
21 0	Mast cells and eosinophils in invasive breast carcinoma. <i>BMC Cancer</i> , 2007 , 7, 165	4.8	57
209	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020 , 52, 56-73	36.3	56
208	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
207	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
206	Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. <i>Genetics in Medicine</i> , 2017 , 19, 599	-60:3	51
205	Relationship of patientsQage to histopathological features of breast tumours in BRCA1 and BRCA2 and mutation-negative breast cancer families. <i>Breast Cancer Research</i> , 2005 , 7, R465-9	8.3	51
204	Effectiveness of a 12-month exercise program on physical performance and quality of life of breast cancer survivors. <i>Anticancer Research</i> , 2012 , 32, 3875-84	2.3	51
203	Molecular subtypes in ductal carcinoma in situ of the breast and their relation to prognosis: a population-based cohort study. <i>BMC Cancer</i> , 2013 , 13, 512	4.8	49
202	Cyclin D1 expression is associated with poor prognostic features in estrogen receptor positive breast cancer. <i>Breast Cancer Research and Treatment</i> , 2009 , 113, 75-82	4.4	49
201	Ten-year follow-up of 3 years of oral adjuvant clodronate therapy shows significant prevention of osteoporosis in early-stage breast cancer. <i>Journal of Clinical Oncology</i> , 2008 , 26, 4289-95	2.2	49
200	Comparison of 6q25 breast cancer hits from Asian and European Genome Wide Association Studies in the Breast Cancer Association Consortium (BCAC). <i>PLoS ONE</i> , 2012 , 7, e42380	3.7	49
199	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2014 , 23, 6096-111	5.6	48

(2015-2001)

198	Survival of breast cancer patients in BRCA1, BRCA2, and non-BRCA1/2 breast cancer families: a relative survival analysis from Finland. <i>International Journal of Cancer</i> , 2001 , 93, 368-72	7.5	48	
197	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	
196	Evaluation of RAD50 in familial breast cancer predisposition. <i>International Journal of Cancer</i> , 2006 , 118, 2911-6	7.5	47	
195	The response evaluation of bone metastases in mammary carcinoma. The value of radiology, scintigraphy, and biochemical markers of bone metabolism. <i>Cancer</i> , 1987 , 60, 2907-12	6.4	47	
194	High expression of cyclin D1 is associated to high proliferation rate and increased risk of mortality in women with ER-positive but not in ER-negative breast cancers. <i>Breast Cancer Research and Treatment</i> , 2017 , 164, 667-678	4.4	46	
193	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019 , 10, 431	17.4	45	
192	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> , 2010 , 102, 650-62	9.7	45	
191	Family history, genetic testing, and clinical risk prediction: pooled analysis of CHEK2 1100delC in 1,828 bilateral breast cancers and 7,030 controls. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009 , 18, 230-4	4	44	
190	TOP2A and HER2 gene amplification as predictors of response to anthracycline treatment in breast cancer. <i>Acta Oncolgica</i> , 2006 , 45, 590-6	3.2	44	
189	Free beta-subunit of human chorionic gonadotropin in serum is a diagnostically sensitive marker of seminomatous testicular cancer. <i>Clinical Chemistry</i> , 2008 , 54, 1840-3	5.5	41	
188	Breast tumors from CHEK2 1100delC-mutation carriers: genomic landscape and clinical implications. <i>Breast Cancer Research</i> , 2011 , 13, R90	8.3	40	
187	Identification of novel genetic markers of breast cancer survival. <i>Journal of the National Cancer Institute</i> , 2015 , 107,	9.7	38	
186	Genetic predisposition to in situ and invasive lobular carcinoma of the breast. <i>PLoS Genetics</i> , 2014 , 10, e1004285	6	38	
185	Clinical importance of genomic imbalances in synovial sarcoma evaluated by comparative genomic hybridization. <i>Cancer Genetics and Cytogenetics</i> , 1999 , 115, 39-46		38	
184	Annexin A1 expression in a pooled breast cancer series: association with tumor subtypes and prognosis. <i>BMC Medicine</i> , 2015 , 13, 156	11.4	37	
183	MicroRNA related polymorphisms and breast cancer risk. <i>PLoS ONE</i> , 2014 , 9, e109973	3.7	37	
182	A cytogenetic study of malignant fibrous histiocytoma. <i>Cancer Genetics and Cytogenetics</i> , 1995 , 85, 91-6		37	
181	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , 2015 , 24, 2966-84	5.6	36	

180	Identification of inherited genetic variations influencing prognosis in early-onset breast cancer. <i>Cancer Research</i> , 2013 , 73, 1883-91	10.1	36
179	Multi-variant pathway association analysis reveals the importance of genetic determinants of estrogen metabolism in breast and endometrial cancer susceptibility. <i>PLoS Genetics</i> , 2010 , 6, e1001012	6	36
178	BARD1 variants Cys557Ser and Val507Met in breast cancer predisposition. <i>European Journal of Human Genetics</i> , 2006 , 14, 167-72	5.3	36
177	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, 285-98	5.6	35
176	A Finnish founder mutation in RAD51D: analysis in breast, ovarian, prostate, and colorectal cancer. Journal of Medical Genetics, 2012 , 49, 429-32	5.8	35
175	Variants on the promoter region of PTEN affect breast cancer progression and patient survival. Breast Cancer Research, 2011 , 13, R130	8.3	34
174	Long-term survival of women with basal-like ductal carcinoma in situ of the breast: a population-based cohort study. <i>BMC Cancer</i> , 2010 , 10, 653	4.8	34
173	Short-term intermittent intravenous clodronate in the prevention of bone loss related to chemotherapy-induced ovarian failure. <i>Breast Cancer Research and Treatment</i> , 2004 , 87, 181-8	4.4	34
172	11q13 is a susceptibility locus for hormone receptor positive breast cancer. <i>Human Mutation</i> , 2012 , 33, 1123-32	4.7	33
171	Complexity of 12q13-22 amplicon in liposarcoma: microsatellite repeat analysis. <i>Genes Chromosomes and Cancer</i> , 1997 , 18, 66-70	5	33
170	The prognostic role of HER2 expression in ductal breast carcinoma in situ (DCIS); a population-based cohort study. <i>BMC Cancer</i> , 2015 , 15, 468	4.8	32
169	A genome-wide association scan on estrogen receptor-negative breast cancer. <i>Breast Cancer Research</i> , 2010 , 12, R93	8.3	32
168	Breast cancer on the Internet: the quality of Swedish breast cancer websites. <i>Breast</i> , 2004 , 13, 376-82	3.6	32
167	Pulmonary toxicity after radiotherapy in primary breast cancer patients: results from a randomized chemotherapy study. <i>International Journal of Radiation Oncology Biology Physics</i> , 2002 , 52, 128-36	4	32
166	Oral trofosfamide: an active drug in the treatment of soft-tissue sarcoma. <i>Cancer Chemotherapy and Pharmacology</i> , 1995 , 36, 263-5	3.5	32
165	Missense variants in ATM in 26,101 breast cancer cases and 29,842 controls. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010 , 19, 2143-51	4	31
164	Combined effects of single nucleotide polymorphisms TP53 R72P and MDM2 SNP309, and p53 expression on survival of breast cancer patients. <i>Breast Cancer Research</i> , 2009 , 11, R89	8.3	31
163	Soft-tissue sarcomas of the upper extremity: surgical treatment and outcome. <i>Plastic and Reconstructive Surgery</i> , 2004 , 113, 222-30; discussion 231-2	2.7	31

(2007-1999)

162	Post-mastectomy radiotherapy in pT3N0M0 breast cancer: is it needed?. <i>Radiotherapy and Oncology</i> , 1999 , 52, 213-7	5.3	31	
161	Genetic predisposition to ductal carcinoma in situ of the breast. <i>Breast Cancer Research</i> , 2016 , 18, 22	8.3	31	
160	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, 934-45	4	29	
159	The relationship between radiation doses to coronary arteries and location of coronary stenosis requiring intervention in breast cancer survivors. <i>Radiation Oncology</i> , 2019 , 14, 40	4.2	28	
158	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	
157	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. <i>Human Molecular Genetics</i> , 2014 , 23, 1934-46	5.6	28	
156	Clinical significance of genetic imbalances revealed by comparative genomic hybridization in chondrosarcomas. <i>Human Pathology</i> , 1999 , 30, 1247-53	3.7	28	
155	Radiation dose distribution in coronary arteries in breast cancer radiotherapy. <i>Acta Oncolgica</i> , 2016 , 55, 959-63	3.2	27	
154	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017 , 46, 1814-1822	7.8	27	
153	Genome-wide scanning for linkage in Finnish breast cancer families. <i>European Journal of Human Genetics</i> , 2004 , 12, 98-104	5.3	27	
152	Radiotherapy in desmoid tumors: Treatment response, local control, and analysis of local failures. <i>Strahlentherapie Und Onkologie</i> , 2017 , 193, 269-275	4.3	26	
151	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	
150	Combined effect of CCND1 and COMT polymorphisms and increased breast cancer risk. <i>BMC Cancer</i> , 2008 , 8, 6	4.8	26	
149	BACH1 Ser919Pro variant and breast cancer risk. <i>BMC Cancer</i> , 2006 , 6, 19	4.8	26	
148	Prophylactic filgrastim (G-CSF) during mitomycin-C, mitoxantrone, and methotrexate (MMM) treatment for metastatic breast cancer. A randomized study. <i>American Journal of Clinical Oncology: Cancer Clinical Trials</i> , 1996 , 19, 232-4	2.7	26	
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	8.3	26	
146	Eukaryotic translation initiation factor 4E (eIF4E) expression is associated with breast cancer tumor phenotype and predicts survival after anthracycline chemotherapy treatment. <i>Breast Cancer Research and Treatment</i> , 2013 , 141, 79-88	4.4	25	
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80 79 78 77	Cytogenetic study of extraskeletal mesenchymal chondrosarcoma. A case report. <i>Cancer Genetics and Cytogenetics</i> , 1996 , 86, 170-3 FEC (5-fluorouracil-epirubicin-cyclophosphamide) monthly versus FEC weekly in metastatic breast cancer. First results of a randomized trial. <i>Acta Oncolgica</i> , 1992 , 31, 231-6 Long-term results of surgical resection of lung metastases from soft tissue sarcoma: A single center experience. <i>Journal of Surgical Oncology</i> , 2019 , 120, 168-175 Recurrent moderate-risk mutations in Finnish breast and ovarian cancer patients. <i>International Journal of Cancer</i> , 2019 , 145, 2692-2700 Transcriptome-wide association study of breast cancer risk by estrogen-receptor status. <i>Genetic</i>	2.8 7·5	10 10 9
80 79 78 77 76	Cytogenetic study of extraskeletal mesenchymal chondrosarcoma. A case report. <i>Cancer Genetics and Cytogenetics</i> , 1996 , 86, 170-3 FEC (5-fluorouracil-epirubicin-cyclophosphamide) monthly versus FEC weekly in metastatic breast cancer. First results of a randomized trial. <i>Acta Oncolgica</i> , 1992 , 31, 231-6 Long-term results of surgical resection of lung metastases from soft tissue sarcoma: A single center experience. <i>Journal of Surgical Oncology</i> , 2019 , 120, 168-175 Recurrent moderate-risk mutations in Finnish breast and ovarian cancer patients. <i>International Journal of Cancer</i> , 2019 , 145, 2692-2700 Transcriptome-wide association study of breast cancer risk by estrogen-receptor status. <i>Genetic Epidemiology</i> , 2020 , 44, 442-468 Germline variation in TP53 regulatory network genes associates with breast cancer survival and	2.8 7·5 2.6	10 10 9 9 9

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17	Common variants in breast cancer risk loci predispose to distinct tumor subtypes		1
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15	Risk of primary lung cancer after adjuvant radiotherapy in breast cancer-a large population-based study. <i>Npj Breast Cancer</i> , 2021 , 7, 71	7.8	1
14	Further validation of the Toronto extremity salvage score for lower extremity soft tissue sarcoma based on Finnish patients. <i>Journal of Plastic, Reconstructive and Aesthetic Surgery</i> , 2021 , 74, 71-78	1.7	1
13	Sense of Coherence as Predictor of Quality of Life in Early Breast Cancer Patients. <i>Anticancer Research</i> , 2021 , 41, 5045-5052	2.3	1
12	Performance of automated scoring of ER, PR, HER2, CK5/6 and EGFR in breast cancer tissue microarrays in the Breast Cancer Association Consortium 2014 , n/a-n/a		1
11	A graphical LASSO analysis of global quality of life, sub scales of the EORTC QLQ-C30 instrument and depression in early breast cancer <i>Scientific Reports</i> , 2022 , 12, 2112	4.9	O
10	Single centre 30-year experience in treating retroperitoneal liposarcomas. <i>Journal of Surgical Oncology</i> , 2020 , 122, 1163-1172	2.8	O
9	Monitoring serum estradiol levels in breast cancer patients during extended adjuvant letrozole treatment after five years of tamoxifen: a prospective trial. <i>Breast Cancer Research and Treatment</i> , 2021 , 187, 769-775	4.4	O
8	Expression of markers of stem cell characteristics, epithelial-mesenchymal transition, basal-like phenotype, proliferation, and androgen receptor in metaplastic breast cancer and their prognostic impact. <i>Acta Oncolgica</i> , 2021 , 60, 1233-1239	3.2	О
7	A search for modifying genetic factors in CHEK2:c.1100delC breast cancer patients. <i>Scientific Reports</i> , 2021 , 11, 14763	4.9	O
6	Reply-Letter to the editor. <i>Breast Journal</i> , 2019 , 25, 1332	1.2	
5	Large-Scale Genomic Analyses Link Reproductive Aging to Hypothalamic Signaling, Breast Cancer Susceptibility, and BRCA1-Mediated DNA Repair. <i>Obstetrical and Gynecological Survey</i> , 2015 , 70, 758-76	52 ^{2.4}	
4	Breast cancer radiotherapy and coronary artery disease: hazards and protection of organs at risk. Breast Cancer Management, 2012 , 1, 13-16	0.7	
3	Results of treatment in testicular nonseminoma. <i>Annals of Medicine</i> , 1996 , 28, 311-4	1.5	
2	Abstract P3-20-02: The association of clinicopathological variables and patient is preference with surgical decision-making for early breast cancer. <i>Cancer Research</i> , 2022 , 82, P3-20-02-P3-20-02	10.1	
1	Serum Concentration of Thymidine Kinase 1 (TK1) as a Tumor Marker in Soft Tissue Sarcomas <i>Anticancer Research</i> , 2022 , 42, 1509-1515	2.3	