Kari Hemminki

List of Publications by Year
in descending order

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600	29,109	9264 74	9103 144
papers	citations	h-index	g-index

607	607	607	31684
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Incidence trends in lung and bladder cancers in the Nordic Countries before and after the smoking epidemic. European Journal of Cancer Prevention, 2022, 31, 228-234.	1.3	9
2	Survival in bladder and upper urinary tract cancers in Finland and Sweden through 50 years. PLoS ONE, 2022, 17, e0261124.	2.5	4
3	T- and B-Cells in the Inner Invasive Margin of Hepatocellular Carcinoma after Resection Associate with Favorable Prognosis. Cancers, 2022, 14, 604.	3.7	4
4	Whole-Exome Sequencing Identifies a Novel Germline Variant in PTK7 Gene in Familial Colorectal Cancer. International Journal of Molecular Sciences, 2022, 23, 1295.	4.1	2
5	Functional dissection of inherited non-coding variation influencing multiple myeloma risk. Nature Communications, 2022, 13, 151.	12.8	10
6	Germline Variants of CYBA and TRPM4 Predispose to Familial Colorectal Cancer. Cancers, 2022, 14, 670.	3.7	11
7	Longâ€term incidence and survival trends in cancer of the gallbladder and extrahepatic bile ducts in Denmark, Finland, Norway and Sweden with etiological implications related to Thorotrast. International Journal of Cancer, 2022, 151, 200-208.	5.1	7
8	Incidence and survival in oral and pharyngeal cancers in Finland and Sweden through half century. BMC Cancer, 2022, 22, 227.	2.6	6
9	Familial Risks for Liver, Gallbladder and Bile Duct Cancers and for Their Risk Factors in Sweden, a Low-Incidence Country. Cancers, 2022, 14, 1938.	3.7	5
10	Genome-wide meta-analysis of monoclonal gammopathy of undetermined significance (MGUS) identifies risk loci impacting IRF-6. Blood Cancer Journal, 2022, 12, 60.	6.2	2
11	Longâ€term incidence in hepatocellular carcinoma and intrahepatic bile duct cancer in Denmark, Finland, Norway and Sweden, role of Thorotrast?. International Journal of Cancer, 2022, 151, 510-517.	5.1	4
12	Cervical, vaginal and vulvar cancer incidence and survival trends in Denmark, Finland, Norway and Sweden with implications to treatment. BMC Cancer, 2022, 22, 456.	2.6	9
13	Whole exome sequencing identifies novel germline variants of SLC15A4 gene as potentially cancer predisposing in familial colorectal cancer. Molecular Genetics and Genomics, 2022, , 1.	2.1	1
14	Validation and functional characterization of GWAS-identified variants for chronic lymphocytic leukemia: a CRuCIAL study. Blood Cancer Journal, 2022, 12, 79.	6.2	1
15	Incidence and survival in laryngeal and lung cancers in Finland and Sweden through a half century. PLoS ONE, 2022, 17, e0268922.	2.5	8
16	Whole Exome Sequencing Identifies APCDD1 and HDAC5 Genes as Potentially Cancer Predisposing in Familial Colorectal Cancer. International Journal of Molecular Sciences, 2021, 22, 1837.	4.1	6
17	Characterization of rare germline variants in familial multiple myeloma. Blood Cancer Journal, 2021, 11, 33.	6.2	7
18	Second Primary Cancers After Kidney Cancers, and Kidney Cancers as Second Primary Cancers. European Urology Open Science, 2021, 24, 52-59.	0.4	0

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19	Whole Genome Sequencing Prioritizes CHEK2, EWSR1, and TIAM1 as Possible Predisposition Genes for Familial Non-Medullary Thyroid Cancer. Frontiers in Endocrinology, 2021, 12, 600682.	3.5	13
20	Polymorphisms within Autophagy-Related Genes Influence the Risk of Developing Colorectal Cancer: A Meta-Analysis of Four Large Cohorts. Cancers, 2021, 13, 1258.	3.7	3
21	A Novel Low-Risk Germline Variant in the SH2 Domain of the SRC Gene Affects Multiple Pathways in Familial Colorectal Cancer. Journal of Personalized Medicine, 2021, 11, 262.	2.5	0
22	DNA repair gene polymorphisms and chromosomal aberrations in healthy, nonsmoking population. DNA Repair, 2021, 101, 103079.	2.8	3
23	Incidence trends in bladder and lung cancers between Denmark, Finland and Sweden may implicate oral tobacco (snuff/snus) as a possible risk factor. BMC Cancer, 2021, 21, 604.	2.6	7
24	Family history of any cancer for childhood leukemia patients in Sweden. EJHaem, 2021, 2, 421-427.	1.0	1
25	Progress in survival in renal cell carcinoma through 50 years evaluated in Finland and Sweden. PLoS ONE, 2021, 16, e0253236.	2.5	13
26	Bladder and upper urinary tract cancers as first and second primary cancers. Cancer Reports, 2021, 4, e1406.	1.4	7
27	DNA Repair Gene Polymorphisms and Chromosomal Aberrations in Exposed Populations. Frontiers in Genetics, 2021, 12, 691947.	2.3	3
28	Family history of early onset acute lymphoblastic leukemia is suggesting genetic associations. Scientific Reports, 2021, 11, 12370.	3.3	2
29	Search for AL amyloidosis risk factors using Mendelian randomization. Blood Advances, 2021, 5, 2725-2731.	5.2	5
30	Second Primary Cancers After Gastric Cancer, and Gastric Cancer as Second Primary Cancer. Clinical Epidemiology, 2021, Volume 13, 515-525.	3.0	9
31	Combinations of Low-Frequency Genetic Variants Might Predispose to Familial Pancreatic Cancer. Journal of Personalized Medicine, 2021, 11, 631.	2.5	9
32	Survival in colon and rectal cancers in Finland and Sweden through 50 years. BMJ Open Gastroenterology, 2021, 8, e000644.	2.7	16
33	Epidemiology of Amyloidosis and Genetic Pathways to Diagnosis and Typing. Hemato, 2021, 2, 429-440.	0.6	0
34	Second Primary Cancers After Liver, Gallbladder and Bile Duct Cancers, and These Cancers as Second Primary Cancers. Clinical Epidemiology, 2021, Volume 13, 683-691.	3.0	6
35	Incidence, mortality and survival in multiple myeloma compared to other hematopoietic neoplasms in Sweden up to year 2016. Scientific Reports, 2021, 11, 17272.	3.3	12
36	Epidemiology, genetics and treatment of multiple myeloma and precursor diseases. International Journal of Cancer, 2021, 149, 1980-1996.	5.1	25

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37	Familial Risks and Proportions Describing Population Landscape of Familial Cancer. Cancers, 2021, 13, 4385.	3.7	20
38	Family History of Head and Neck Cancers. Cancers, 2021, 13, 4115.	3.7	2
39	A rare large duplication of MLH1 identified in Lynch syndrome. Hereditary Cancer in Clinical Practice, 2021, 19, 10.	1.5	2
40	Familial Risks between Pernicious Anemia and Other Autoimmune Diseases in the Population of Sweden. Autoimmune Diseases, 2021, 2021, 1-5.	0.6	4
41	Prevalence of the GFI1-36N SNP in Multiple Myeloma Patients and Its Impact on the Prognosis. Frontiers in Oncology, 2021, 11, 757664.	2.8	3
42	The Asthma Family Tree: Evaluating Associations Between Childhood, Parental, and Grandparental Asthma in Seven Chinese Cities. Frontiers in Pediatrics, 2021, 9, 720273.	1.9	4
43	Types of second primary cancer influence overall survival in cutaneous melanoma. BMC Cancer, 2021, 21, 1123.	2.6	3
44	Incidence, mortality and survival in malignant pleural mesothelioma before and after asbestos in Denmark, Finland, Norway and Sweden. BMC Cancer, 2021, 21, 1189.	2.6	11
45	Second primary cancers in nonâ€Hodgkin lymphoma: Family history and survival. International Journal of Cancer, 2020, 146, 970-976.	5.1	15
46	Autoimmune diseases and hematological malignancies: Exploring the underlying mechanisms from epidemiological evidence. Seminars in Cancer Biology, 2020, 64, 114-121.	9.6	20
47	Genetic epidemiology of colorectal cancer and associated cancers. Mutagenesis, 2020, 35, 207-219.	2.6	17
48	Second Primary Cancers in Patients with Invasive and In Situ Squamous Cell Skin Carcinoma, Kaposi Sarcoma, and Merkel Cell Carcinoma: Role for Immune Mechanisms?. Journal of Investigative Dermatology, 2020, 140, 48-55.e1.	0.7	7
49	Eight novel loci implicate shared genetic etiology in multiple myeloma, AL amyloidosis, and monoclonal gammopathy of unknown significance. Leukemia, 2020, 34, 1187-1191.	7.2	13
50	Genetic Variants Associated with Chronic Kidney Disease in a Spanish Population. Scientific Reports, 2020, 10, 144.	3.3	29
51	Genetic predisposition for multiple myeloma. Leukemia, 2020, 34, 697-708.	7.2	25
52	Epistatic effect of TLR3 and cGASâ€STINGâ€IKKÎµâ€TBK1â€IFN signaling variants on colorectal cancer risk. Cancer Medicine, 2020, 9, 1473-1484.	2.8	10
53	Genome-wide study on uveal melanoma patients finds association to DNA repair gene TDP1. Melanoma Research, 2020, 30, 166-172.	1.2	6
54	Familial associations for rheumatoid autoimmune diseases. Rheumatology Advances in Practice, 2020, 4, rkaa048.	0.7	7

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55	Cancer Predisposition Genes in Cancer-Free Families. Cancers, 2020, 12, 2770.	3.7	2
56	Familial risks between Graves disease and Hashimoto thyroiditis and other autoimmune diseases in the population of Sweden. Journal of Translational Autoimmunity, 2020, 3, 100058.	4.0	20
57	Informing patients about their mutation tests: CDKN2A c.256G>A in melanoma as an example. Hereditary Cancer in Clinical Practice, 2020, 18, 15.	1.5	3
58	Rate differences between first and second primary cancers may outline immune dysfunction as a key risk factor. Cancer Medicine, 2020, 9, 8258-8265.	2.8	9
59	Incidence Differences Between First Primary Cancers and Second Primary Cancers Following Skin Squamous Cell Carcinoma as Etiological Clues . Clinical Epidemiology, 2020, Volume 12, 857-864.	3.0	4
60	Genomic imprinting analyses identify maternal effects as a cause of phenotypic variability in type 1 diabetes and rheumatoid arthritis. Scientific Reports, 2020, 10, 11562.	3.3	11
61	Familial risks between giant cell arteritis and Takayasu arteritis and other autoimmune diseases in the population of Sweden. Scientific Reports, 2020, 10, 20887.	3.3	2
62	Impact of genetic polymorphisms in kinetochore and spindle assembly genes on chromosomal aberration frequency in healthy humans. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2020, 858-860, 503253.	1.7	2
63	Search for multiple myeloma risk factors using Mendelian randomization. Blood Advances, 2020, 4, 2172-2179.	5.2	27
64	Determining the Appropriate Risk-Adapted Screening Age for Familial Breast Cancer. JAMA Oncology, 2020, 6, 933.	7.1	0
65	A Germline Mutation in the POT1 Gene Is a Candidate for Familial Non-Medullary Thyroid Cancer. Cancers, 2020, 12, 1441.	3.7	24
66	Identification of Familial Hodgkin Lymphoma Predisposing Genes Using Whole Genome Sequencing. Frontiers in Bioengineering and Biotechnology, 2020, 8, 179.	4.1	12
67	Second Primary Cancers in Melanoma Patients Critically Shorten Survival . Clinical Epidemiology, 2020, Volume 12, 105-112.	3.0	3
68	Loci associated with genomic damage levels in chronic kidney disease patients and controls. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2020, 852, 503167.	1.7	10
69	Association between tumor characteristics and second primary cancers with cutaneous melanoma survival: A nationwide cohort study. Pigment Cell and Melanoma Research, 2020, 33, 625-632.	3.3	3
70	TERT promoter mutations in actinic keratosis before and after treatment. International Journal of Cancer, 2020, 146, 2932-2934.	5.1	6
71	Telomere length in peripheral blood lymphocytes related to genetic variation in telomerase, prognosis and clinicopathological features in breast cancer patients. Mutagenesis, 2020, 35, 491-497.	2.6	11
72	Familial associations between autoimmune hepatitis and primary biliary cholangitis and other autoimmune diseases. PLoS ONE, 2020, 15, e0240794.	2.5	3

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73	Chromosomal damage and telomere length in peripheral blood lymphocytes of cancer patients. Oncology Reports, 2020, 44, 2219-2230.	2.6	4
74	Familial associations for Addison's disease and between Addison's disease and other autoimmune diseases. Endocrine Connections, 2020, 9, 1114-1120.	1.9	0
75	Familial associations for Addisonâ€™s disease and between Addisonâ€™s disease and other autoimmune diseases. Endocrine Connections, 2020, 9, 1114-1120.	1.9	2
76	Characterization of Rare Germline Variants in Familial Multiple Myeloma. Blood, 2020, 136, 45-46.	1.4	0
77	Analysis of 153â€‰115 patients with hematological malignancies refines the spectrum of familial risk. Blood, 2019, 134, 960-969.	1.4	51
78	Familial Clustering, Second Primary Cancers and Causes of Death in Penile, Vulvar and Vaginal Cancers. Scientific Reports, 2019, 9, 11804.	3.3	9
79	Familial Cancer: How to Successfully Recruit Families for Germline Mutations Studies? Multiple Myeloma as an Example. Clinical Lymphoma, Myeloma and Leukemia, 2019, 19, 635-644.e2.	0.4	4
80	Whole Genome Sequencing of Familial Non-Medullary Thyroid Cancer Identifies Germline Alterations in MAPK/ERK and PI3K/AKT Signaling Pathways. Biomolecules, 2019, 9, 605.	4.0	27
81	Comparison of Familial Clustering of Anogenital and Skin Cancers Between In Situ and Invasive Types. Scientific Reports, 2019, 9, 16151.	3.3	2
82	Transcriptome-wide association study of multiple myeloma identifies candidate susceptibility genes. Human Genomics, 2019, 13, 37.	2.9	14
83	Distinct pathways associated with chromosomal aberration frequency in a cohort exposed to genotoxic compounds compared to general population. Mutagenesis, 2019, 34, 323-330.	2.6	6
84	Familial Associations of Colon and Rectal Cancers With Other Cancers. Diseases of the Colon and Rectum, 2019, 62, 189-195.	1.3	5
85	Second primary cancers in patients with acute lymphoblastic, chronic lymphocytic and hairy cell leukaemia. British Journal of Haematology, 2019, 185, 232-239.	2.5	34
86	Types of second primary cancers influence survival in chronic lymphocytic and hairy cell leukemia patients. Blood Cancer Journal, 2019, 9, 40.	6.2	7
87	Single nucleotide polymorphisms within MUC4 are associated with colorectal cancer survival. PLoS ONE, 2019, 14, e0216666.	2.5	15
88	Update on genetic predisposition to colorectal cancer and polyposis. Molecular Aspects of Medicine, 2019, 69, 10-26.	6.4	113
89	Genome-wide interaction and pathway-based identification of key regulators in multiple myeloma. Communications Biology, 2019, 2, 89.	4.4	14
90	Second cancers and causes of death in patients with testicular cancer in Sweden. PLoS ONE, 2019, 14, e0214410.	2.5	15

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91	Genome-wide association study of monoclonal gammopathy of unknown significance (MGUS): comparison with multiple myeloma. Leukemia, 2019, 33, 1817-1821.	7.2	14
92	Associations between autoimmune conditions and hepatobiliary cancer risk among elderly US adults. International Journal of Cancer, 2019, 144, 707-717.	5.1	20
93	Impact of family history of cancer on risk and mortality of second cancers in patients with prostate cancer. Prostate Cancer and Prostatic Diseases, 2019, 22, 143-149.	3.9	12
94	Second primary cancer after female breast cancer: Familial risks and cause of death. Cancer Medicine, 2019, 8, 400-407.	2.8	13
95	Genetic correlation between multiple myeloma and chronic lymphocytic leukaemia provides evidence for shared aetiology. Blood Cancer Journal, 2019, 9, 1.	6.2	40
96	Genetic variation associated with chromosomal aberration frequency: A genomeâ€wide association study. Environmental and Molecular Mutagenesis, 2019, 60, 17-28.	2.2	9
97	Familial Risks Between Urolithiasis and Cancer. Scientific Reports, 2018, 8, 3083.	3.3	1
98	Cytogenetic aberrations in multiple myeloma are associated with shifts in serum immunoglobulin isotypes distribution and levels. Haematologica, 2018, 103, e162-e164.	3.5	5
99	RE: Familial Cancer Clustering of Urothelial Cancer: A Population-Based Caseâ€“Control Study. Journal of the National Cancer Institute, 2018, 110, 1277-1278.	6.3	1
100	Genome-wide association study identifies susceptibility loci for B-cell childhood acute lymphoblastic leukemia. Nature Communications, 2018, 9, 1340.	12.8	58
101	The multiple myeloma risk allele at 5q15 lowers ELL2 expression and increases ribosomal gene expression. Nature Communications, 2018, 9, 1649.	12.8	22
102	Whole genome sequencing reveals DICER1 as a candidate predisposing gene in familial Hodgkin lymphoma. International Journal of Cancer, 2018, 143, 2076-2078.	5.1	19
103	Bortezomibâ€induced peripheral neuropathy: A genomeâ€wide association study on multiple myeloma patients. Hematological Oncology, 2018, 36, 232-237.	1.7	20
104	Bleomycinâ€induced chromosomal damage and shortening of telomeres in peripheral blood lymphocytes of incident cancer patients. Genes Chromosomes and Cancer, 2018, 57, 61-69.	2.8	12
105	Familial risks in urolithiasis in the population of Sweden. BJU International, 2018, 121, 479-485.	2.5	18
106	Response: Methods for second primary cancers evaluation have to be standardized. International Journal of Cancer, 2018, 142, 1286-1287.	5.1	0
107	Familial Risks and Mortality in Second Primary Cancers in Melanoma. JNCI Cancer Spectrum, 2018, 2, pky068.	2.9	12
108	Familial risks of ovarian cancer by age at diagnosis, proband type and histology. PLoS ONE, 2018, 13, e0205000.	2.5	22

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109	Importance of tumor location and histology in familial risk of upper gastrointestinal cancers: a nationwide cohort study. Clinical Epidemiology, 2018, Volume 10, 1169-1179.	3.0	10
110	Prostate cancer survivors: Risk and mortality in second primary cancers. Cancer Medicine, 2018, 7, 5752-5759.	2.8	17
111	Second primary cancers in nonâ€Hodgkin lymphoma: Bidirectional analyses suggesting role for immune dysfunction. International Journal of Cancer, 2018, 143, 2449-2457.	5.1	22
112	Familial risks of second primary cancers and mortality in ovarian cancer patients. Clinical Epidemiology, 2018, Volume 10, 1457-1466.	3.0	10
113	Clinical landscape of cancer metastases. Cancer Medicine, 2018, 7, 5534-5542.	2.8	74
114	Identification of multiple risk loci and regulatory mechanisms influencing susceptibility to multiple myeloma. Nature Communications, 2018, 9, 3707.	12.8	86
115	Genome-wide association study implicates immune dysfunction in the development of Hodgkin lymphoma. Blood, 2018, 132, 2040-2052.	1.4	17
116	Enrichment of B cell receptor signaling and epidermal growth factor receptor pathways in monoclonal gammopathy of undetermined significance: a genome-wide genetic interaction study. Molecular Medicine, 2018, 24, 30.	4.4	9
117	Familial risk of pleural mesothelioma increased drastically in certain occupations: A nationwide prospective cohort study. European Journal of Cancer, 2018, 103, 1-6.	2.8	12
118	Short article: Influence of regulatory NLRC5 variants on colorectal cancer survival and 5-fluorouracil-based chemotherapy. European Journal of Gastroenterology and Hepatology, 2018, 30, 838-842.	1.6	6
119	Genetic variation of acquired structural chromosomal aberrations. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2018, 836, 13-21.	1.7	19
120	Familial Urinary Bladder Cancer with Other Cancers. European Urology Oncology, 2018, 1, 461-466.	5.4	4
121	Coding variants in NOD-like receptors: An association study on risk and survival of colorectal cancer. PLoS ONE, 2018, 13, e0199350.	2.5	6
122	Risk of second primary cancer following myeloid neoplasia and risk of myeloid neoplasia as second primary cancer: a nationwide, observational follow up study in Sweden. Lancet Haematology,the, 2018, 5, e368-e377.	4.6	14
123	Familial risks in and between stone diseases: sialolithiasis, urolithiasis and cholelithiasis in the population of Sweden. BMC Nephrology, 2018, 19, 158.	1.8	5
124	Familial risks of acute myeloid leukemia, myelodysplastic syndromes, and myeloproliferative neoplasms. Blood, 2018, 132, 973-976.	1.4	35
125	Borderline Ovarian Tumors Share Familial Risks with Themselves and Invasive Cancers. Cancer Epidemiology Biomarkers and Prevention, 2018, 27, 1358-1363.	2.5	2
126	Familial Associations in Testicular Cancer with Other Cancers. Scientific Reports, 2018, 8, 10880.	3.3	12

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127	Levels of DNA damage (Micronuclei) in patients suffering from chronic kidney disease. Role of GST polymorphisms. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2018, 836, 41-46.	1.7	6
128	HLAandKIRAssociations of Cervical Neoplasia. Journal of Infectious Diseases, 2018, 218, 2006-2015.	4.0	22
129	Chemotherapy-induced peripheral neuropathy: evidence from genome-wide association studies and replication within multiple myeloma patients. BMC Cancer, 2018, 18, 820.	2.6	18
130	Familial Ovarian Cancer ClustersÂwith Other Cancers. Scientific Reports, 2018, 8, 11561.	3.3	6
131	Multiple myeloma: family history and mortality in second primary cancers. Blood Cancer Journal, 2018, 8, 75.	6.2	5
132	Familial Cancer Variant Prioritization Pipeline version 2 (FCVPPv2) applied to a papillary thyroid cancer family. Scientific Reports, 2018, 8, 11635.	3.3	30
133	Investigation of single and synergic effects of NLRC5 and PD-L1 variants on the risk of colorectal cancer. PLoS ONE, 2018, 13, e0192385.	2.5	20
134	Single nucleotide polymorphisms within Mucin-type O-glycan genes are associated with colorectal cancer survival.. Journal of Clinical Oncology, 2018, 36, e15607-e15607.	1.6	0
135	Genome-wide association analysis of chronic lymphocytic leukaemia, Hodgkin lymphoma and multiple myeloma identifies pleiotropic risk loci. Scientific Reports, 2017, 7, 41071.	3.3	31
136	Risk of other Cancers in Families with Melanoma: Novel Familial Links. Scientific Reports, 2017, 7, 42601.	3.3	23
137	Common cancers share familial susceptibility: implications for cancer genetics and counselling. Journal of Medical Genetics, 2017, 54, 248-253.	3.2	12
138	Functional germline variants in driver genes of breast cancer. Cancer Causes and Control, 2017, 28, 259-271.	1.8	12
139	Low expression of hexokinase-2 is associated with false-negative FDGâ€“positron emission tomography in multiple myeloma. Blood, 2017, 130, 30-34.	1.4	180
140	Genetics of gallbladder cancer. Lancet Oncology, The, 2017, 18, e296.	10.7	9
141	Genomewide association study on monoclonal gammopathy of unknown significance (MGUS). European Journal of Haematology, 2017, 99, 70-79.	2.2	16
142	Novel recurrent chromosomal aberrations detected in clonal plasma cells of light chain amyloidosis patients show potential adverse prognostic effect: first results from a genome-wide copy number array analysis. Haematologica, 2017, 102, 1281-1290.	3.5	15
143	Concordant and discordant familial cancer: Familial risks, proportions and population impact. International Journal of Cancer, 2017, 140, 1510-1516.	5.1	57
144	Familial associations of male breast cancer with other cancers. Breast Cancer Research and Treatment, 2017, 166, 897-902.	2.5	7

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145	Genetic Predisposition to Multiple Myeloma at 5q15 Is Mediated by an ELL2 Enhancer Polymorphism. Cell Reports, 2017, 20, 2556-2564.	6.4	17
146	Surveillance Bias in Cancer Risk After Unrelated Medical Conditions: Example Urolithiasis. Scientific Reports, 2017, 7, 8073.	3.3	21
147	Other cancers in lung cancer families are overwhelmingly smoking-related cancers. ERJ Open Research, 2017, 3, 00006-2017.	2.6	3
148	Risk of second primary cancers in women diagnosed with endometrial cancer in G erman and S wedish cancer registries. International Journal of Cancer, 2017, 141, 2270-2280.	5.1	13
149	Familial associations of female breast cancer with other cancers. International Journal of Cancer, 2017, 141, 2253-2259.	5.1	19
150	Genome-wide association study of classical Hodgkin lymphoma identifies key regulators of disease susceptibility. Nature Communications, 2017, 8, 1892.	12.8	40
151	Familial Associations of Colorectal Cancer with Other Cancers. Scientific Reports, 2017, 7, 5243.	3.3	6
152	Genome-wide association study of clinical parameters in immunoglobulin light chain amyloidosis in three patient cohorts. Haematologica, 2017, 102, e411-e414.	3.5	7
153	Whole-exome sequencing identifies novel candidate predisposition genes for familial polycythemia vera. Human Genomics, 2017, 11, 6.	2.9	11
154	Familial Associations Between Prostate Cancer and Other Cancers. European Urology, 2017, 71, 162-165.	1.9	19
155	Identification of miRSNPs associated with the risk of multiple myeloma. International Journal of Cancer, 2017, 140, 526-534.	5.1	8
156	Genetic Susceptibility to Bortezomib-Induced Peripheral Neuroropathy: Replication of the Reported Candidate Susceptibility Loci. Neurochemical Research, 2017, 42, 925-931.	3.3	15
157	Familial risks for gallstones in the population of Sweden. BMJ Open Gastroenterology, 2017, 4, e000188.	2.7	9
158	Direct evidence for a polygenic etiology in familial multiple myeloma. Blood Advances, 2017, 1, 619-623.	5.2	15
159	Risk of Second Cancer in Hodgkin Lymphoma Survivors and Influence of Family History. Journal of Clinical Oncology, 2017, 35, 1584-1590.	1.6	61
160	Defining the genetic susceptibility to cervical neoplasiaâ€"A genome-wide association study. PLoS Genetics, 2017, 13, e1006866.	3.5	105
161	Inherited variants in genes somatically mutated in thyroid cancer. PLoS ONE, 2017, 12, e0174995.	2.5	5
162	Second cancer risk following Hodgkin lymphoma. Oncotarget, 2017, 8, 78261-78262.	1.8	5

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163	Metastatic spread in patients with gastric cancer. Oncotarget, 2016, 7, 52307-52316.	1.8	272
164	Origin of B-Cell Neoplasms in Autoimmune Disease. PLoS ONE, 2016, 11, e0158360.	2.5	17
165	<scp>TERT</scp> promoter mutations in melanoma survival. International Journal of Cancer, 2016, 139, 75-84.	5.1	101
166	Cancer of unknown primary is associated with diabetes. European Journal of Cancer Prevention, 2016, 25, 246-251.	1.3	14
167	Location of metastases in cancer of unknown primary are not random and signal familial clustering. Scientific Reports, 2016, 6, 22891.	3.3	14
168	The Incidence of Senile Cataract and Glaucoma is Increased in Patients with Plasma Cell Dyscrasias: Etiologic Implications. Scientific Reports, 2016, 6, 28500.	3.3	14
169	Predictive and Prognostic Clinical Variables in Cancer Patients Treated With Adenoviral Oncolytic Immunotherapy. Molecular Therapy, 2016, 24, 1323-1332.	8.2	28
170	Analysis of functional germline variants in APOBEC3 and driver genes on breast cancer risk in Moroccan study population. BMC Cancer, 2016, 16, 165.	2.6	20
171	Polymorphisms within base and nucleotide excision repair pathways and risk of differentiated thyroid carcinoma. DNA Repair, 2016, 41, 27-31.	2.8	5
172	Runs of homozygosity and inbreeding in thyroid cancer. BMC Cancer, 2016, 16, 227.	2.6	17
173	Genetic variation in the major mitotic checkpoint genes associated with chromosomal aberrations in healthy humans. Cancer Letters, 2016, 380, 442-446.	7.2	12
174	The epidemiology of metastases in neuroendocrine tumors. International Journal of Cancer, 2016, 139, 2679-2686.	5.1	233
175	Multiple myeloma risk variant at 7p15.3 creates an IRF4-binding site and interferes with CDCA7L expression. Nature Communications, 2016, 7, 13656.	12.8	32
176	Genome-wide association study identifies multiple susceptibility loci for multiple myeloma. Nature Communications, 2016, 7, 12050.	12.8	146
177	Pedigree based DNA sequencing pipeline for germline genomes of cancer families. Hereditary Cancer in Clinical Practice, 2016, 14, 16.	1.5	7
178	Patterns of metastasis in colon and rectal cancer. Scientific Reports, 2016, 6, 29765.	3.3	652
179	Risk of Second Primary Cancers in Multiple Myeloma Survivors in German and Swedish Cancer Registries. Scientific Reports, 2016, 6, 22084.	3.3	15
180	Age-Dependent Metastatic Spread and Survival: Cancer of Unknown Primary as a Model. Scientific Reports, 2016, 6, 23725.	3.3	14

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181	Risk of second primary cancers after malignant mesothelioma and vice versa. Cancer Letters, 2016, 379, 94-99.	7.2	8
182	Survival in familial and non-familial breast cancer by age and stage at diagnosis. European Journal of Cancer, 2016, 52, 10-18.	2.8	14
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