Kari Hemminki

List of Publications by Year in Descending Order

587 papers	23,689 citations	67 h-index	132 g-index
607 ext. papers	26,602 ext. citations	5.8 avg, IF	6.86 L-index

#	Paper	IF	Citations
587	Survival in bladder and upper urinary tract cancers in Finland and Sweden through 50 years.. *PLoS ONE*, **2022**, 17, e0261124	3.7	0
586	Functional dissection of inherited non-coding variation influencing multiple myeloma risk.. *Nature Communications*, **2022**, 13, 151	17.4	0
585	Germline Variants of and Predispose to Familial Colorectal Cancer.. *Cancers*, **2022**, 14,	6.6	3
584	Incidence and survival in oral and pharyngeal cancers in Finland and Sweden through half century.. *BMC Cancer*, **2022**, 22, 227	4.8	1
583	Genome-wide meta-analysis of monoclonal gammopathy of undetermined significance (MGUS) identifies risk loci impacting IRF-6.. *Blood Cancer Journal*, **2022**, 12, 60	7	
582	Cervical, vaginal and vulvar cancer incidence and survival trends in Denmark, Finland, Norway and Sweden with implications to treatment.. *BMC Cancer*, **2022**, 22, 456	4.8	1
581	Whole exome sequencing identifies novel germline variants of SLC15A4 gene as potentially cancer predisposing in familial colorectal cancer.. *Molecular Genetics and Genomics*, **2022**, 1	3.1	
580	Validation and functional characterization of GWAS-identified variants for chronic lymphocytic leukemia: a CRuCIAL study.. *Blood Cancer Journal*, **2022**, 12, 79	7	
579	Incidence and survival in laryngeal and lung cancers in Finland and Sweden through a half century. *PLoS ONE*, **2022**, 17, e0268922	3.7	0
578	Incidence, mortality and survival in malignant pleural mesothelioma before and after asbestos in Denmark, Finland, Norway and Sweden. *BMC Cancer*, **2021**, 21, 1189	4.8	2
577	Prevalence of the GFI1-36N SNP in Multiple Myeloma Patients and Its Impact on the Prognosis. *Frontiers in Oncology*, **2021**, 11, 757664	5.3	0
576	The Asthma Family Tree: Evaluating Associations Between Childhood, Parental, and Grandparental Asthma in Seven Chinese Cities. *Frontiers in Pediatrics*, **2021**, 9, 720273	3.4	1
575	Types of second primary cancer influence overall survival in cutaneous melanoma. *BMC Cancer*, **2021**, 21, 1123	4.8	0
574	DNA repair gene polymorphisms and chromosomal aberrations in healthy, nonsmoking population. *DNA Repair*, **2021**, 101, 103079	4.3	2
573	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	4.8	3
572	Family history of any cancer for childhood leukemia patients in Sweden. *EJHaem*, **2021**, 2, 421-427	0.9	
571	Progress in survival in renal cell carcinoma through 50 years evaluated in Finland and Sweden. *PLoS ONE*, **2021**, 16, e0253236	3.7	5

570	Bladder and upper urinary tract cancers as first and second primary cancers. *Cancer Reports*, **2021**, e1406	1.5	2
569	Incidence trends in lung and bladder cancers in the Nordic Countries before and after the smoking epidemic. *European Journal of Cancer Prevention*, **2021**,	2	4
568	DNA Repair Gene Polymorphisms and Chromosomal Aberrations in Exposed Populations. *Frontiers in Genetics*, **2021**, 12, 691947	4.5	1
567	Family history of early onset acute lymphoblastic leukemia is suggesting genetic associations. *Scientific Reports*, **2021**, 11, 12370	4.9	0
566	Search for AL amyloidosis risk factors using Mendelian randomization. *Blood Advances*, **2021**, 5, 2725-2731	7.8	3
565	Second Primary Cancers After Gastric Cancer, and Gastric Cancer as Second Primary Cancer. *Clinical Epidemiology*, **2021**, 13, 515-525	5.9	1
564	Combinations of Low-Frequency Genetic Variants Might Predispose to Familial Pancreatic Cancer. *Journal of Personalized Medicine*, **2021**, 11,	3.6	2
563	Survival in colon and rectal cancers in Finland and Sweden through 50 years. *BMJ Open Gastroenterology*, **2021**, 8,	3.9	4
562	Whole Exome Sequencing Identifies and Genes as Potentially Cancer Predisposing in Familial Colorectal Cancer. *International Journal of Molecular Sciences*, **2021**, 22,	6.3	2
561	Characterization of rare germline variants in familial multiple myeloma. *Blood Cancer Journal*, **2021**, 11, 33	7	1
560	Second Primary Cancers After Kidney Cancers, and Kidney Cancers as Second Primary Cancers. *European Urology Open Science*, **2021**, 24, 52-59	0.9	
559	Whole Genome Sequencing Prioritizes , and as Possible Predisposition Genes for Familial Non-Medullary Thyroid Cancer. *Frontiers in Endocrinology*, **2021**, 12, 600682	5.7	5
558	Epidemiology of Amyloidosis and Genetic Pathways to Diagnosis and Typing. *Hemato*, **2021**, 2, 429-440	0.2	
557	Second Primary Cancers After Liver, Gallbladder and Bile Duct Cancers, and These Cancers as Second Primary Cancers. *Clinical Epidemiology*, **2021**, 13, 683-691	5.9	2
556	Incidence, mortality and survival in multiple myeloma compared to other hematopoietic neoplasms in Sweden up to year 2016. *Scientific Reports*, **2021**, 11, 17272	4.9	3
555	Epidemiology, genetics and treatment of multiple myeloma and precursor diseases. *International Journal of Cancer*, **2021**, 149, 1980-1996	7.5	5
554	Familial Risks and Proportions Describing Population Landscape of Familial Cancer. *Cancers*, **2021**, 13,	6.6	3
553	Family History of Head and Neck Cancers. *Cancers*, **2021**, 13,	6.6	2

552	A rare large duplication of MLH1 identified in Lynch syndrome. *Hereditary Cancer in Clinical Practice*, **2021**, 19, 10	2.3	1
551	Familial Risks between Pernicious Anemia and Other Autoimmune Diseases in the Population of Sweden. *Autoimmune Diseases*, **2021**, 2021, 8815297	2.9	0
550	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	3	1
549	Search for multiple myeloma risk factors using Mendelian randomization. *Blood Advances*, **2020**, 4, 2172-2179	7.8	11
548	Determining the Appropriate Risk-Adapted Screening Age for Familial Breast Cancer. *JAMA Oncology*, **2020**, 6, 933-934	13.4	
547	A Germline Mutation in the Gene Is a Candidate for Familial Non-Medullary Thyroid Cancer. *Cancers*, **2020**, 12,	6.6	10
546	Identification of Familial Hodgkin Lymphoma Predisposing Genes Using Whole Genome Sequencing. *Frontiers in Bioengineering and Biotechnology*, **2020**, 8, 179	5.8	7
545	Second Primary Cancers in Melanoma Patients Critically Shorten Survival. *Clinical Epidemiology*, **2020**, 12, 105-112	5.9	0
544	Loci associated with genomic damage levels in chronic kidney disease patients and controls. *Mutation Research - Genetic Toxicology and Environmental Mutagenesis*, **2020**, 852, 503167	3	5
543	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	4.5	1
542	TERT promoter mutations in actinic keratosis before and after treatment. *International Journal of Cancer*, **2020**, 146, 2932-2934	7.5	1
541	Familial associations for Addison disease and between Addison disease and other autoimmune diseases. *Endocrine Connections*, **2020**, 9, 1114-1120	3.5	0
540	Characterization of Rare Germline Variants in Familial Multiple Myeloma. *Blood*, **2020**, 136, 45-46	2.2	
539	Familial associations between autoimmune hepatitis and primary biliary cholangitis and other autoimmune diseases. *PLoS ONE*, **2020**, 15, e0240794	3.7	1
538	Familial associations for Addison's disease and between Addison's disease and other autoimmune diseases. *Endocrine Connections*, **2020**, 9, 1114-1120	3.5	
537	Chromosomal damage and telomere length in peripheral blood lymphocytes of cancer patients. *Oncology Reports*, **2020**, 44, 2219-2230	3.5	0
536	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	2.8	2
535	Eight novel loci implicate shared genetic etiology in multiple myeloma, AL amyloidosis, and monoclonal gammopathy of unknown significance. *Leukemia*, **2020**, 34, 1187-1191	10.7	12

534	Genetic Variants Associated with Chronic Kidney Disease in a Spanish Population. *Scientific Reports*, **2020**, 10, 144	4.9	16
533	Genetic predisposition for multiple myeloma. *Leukemia*, **2020**, 34, 697-708	10.7	11
532	Epistatic effect of TLR3 and cGAS-STING-IKK□TBK1-IFN signaling variants on colorectal cancer risk. *Cancer Medicine*, **2020**, 9, 1473-1484	4.8	6
531	Genome-wide study on uveal melanoma patients finds association to DNA repair gene TDP1. *Melanoma Research*, **2020**, 30, 166-172	3.3	2
530	Familial associations for rheumatoid autoimmune diseases. *Rheumatology Advances in Practice*, **2020**, 4, rkaa048	1.1	2
529	Familial risks between Graves disease and Hashimoto thyroiditis and other autoimmune diseases in the population of Sweden. *Journal of Translational Autoimmunity*, **2020**, 3, 100058	4.1	11
528	Informing patients about their mutation tests: CDKN2A c.256G>A in melanoma as an example. *Hereditary Cancer in Clinical Practice*, **2020**, 18, 15	2.3	0
527	Rate differences between first and second primary cancers may outline immune dysfunction as a key risk factor. *Cancer Medicine*, **2020**, 9, 8258-8265	4.8	6
526	Incidence Differences Between First Primary Cancers and Second Primary Cancers Following Skin Squamous Cell Carcinoma as Etiological Clues. *Clinical Epidemiology*, **2020**, 12, 857-864	5.9	3
525	Genomic imprinting analyses identify maternal effects as a cause of phenotypic variability in type 1 diabetes and rheumatoid arthritis. *Scientific Reports*, **2020**, 10, 11562	4.9	6
524	Familial risks between giant cell arteritis and Takayasu arteritis and other autoimmune diseases in the population of Sweden. *Scientific Reports*, **2020**, 10, 20887	4.9	1
523	Second primary cancers in non-Hodgkin lymphoma: Family history and survival. *International Journal of Cancer*, **2020**, 146, 970-976	7.5	5
522	Autoimmune diseases and hematological malignancies: Exploring the underlying mechanisms from epidemiological evidence. *Seminars in Cancer Biology*, **2020**, 64, 114-121	12.7	10
521	Genetic epidemiology of colorectal cancer and associated cancers. *Mutagenesis*, **2020**, 35, 207-219	2.8	5
520	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	4.3	7
519	Transcriptome-wide association study of multiple myeloma identifies candidate susceptibility genes. *Human Genomics*, **2019**, 13, 37	6.8	5
518	Distinct pathways associated with chromosomal aberration frequency in a cohort exposed to genotoxic compounds compared to general population. *Mutagenesis*, **2019**, 34, 323-330	2.8	5
517	Familial Associations of Colon and Rectal Cancers With Other Cancers. *Diseases of the Colon and Rectum*, **2019**, 62, 189-195	3.1	4

516	Second primary cancers in patients with acute lymphoblastic, chronic lymphocytic and hairy cell leukaemia. *British Journal of Haematology*, **2019**, 185, 232-239	4.5	25
515	Types of second primary cancers influence survival in chronic lymphocytic and hairy cell leukemia patients. *Blood Cancer Journal*, **2019**, 9, 40	7	6
514	Single nucleotide polymorphisms within MUC4 are associated with colorectal cancer survival. *PLoS ONE*, **2019**, 14, e0216666	3.7	9
513	Update on genetic predisposition to colorectal cancer and polyposis. *Molecular Aspects of Medicine*, **2019**, 69, 10-26	16.7	59
512	Genome-wide interaction and pathway-based identification of key regulators in multiple myeloma. *Communications Biology*, **2019**, 2, 89	6.7	11
511	Second cancers and causes of death in patients with testicular cancer in Sweden. *PLoS ONE*, **2019**, 14, e0214410	3.7	9
510	Genome-wide association study of monoclonal gammopathy of unknown significance (MGUS): comparison with multiple myeloma. *Leukemia*, **2019**, 33, 1817-1821	10.7	12
509	Analysis of 153 115 patients with hematological malignancies refines the spectrum of familial risk. *Blood*, **2019**, 134, 960-969	2.2	29
508	Familial Clustering, Second Primary Cancers and Causes of Death in Penile, Vulvar and Vaginal Cancers. *Scientific Reports*, **2019**, 9, 11804	4.9	5
507	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	2	1
506	Whole Genome Sequencing of Familial Non-Medullary Thyroid Cancer Identifies Germline Alterations in MAPK/ERK and PI3K/AKT Signaling Pathways. *Biomolecules*, **2019**, 9,	5.9	18
505	Comparison of Familial Clustering of Anogenital and Skin Cancers Between In Situ and Invasive Types. *Scientific Reports*, **2019**, 9, 16151	4.9	2
504	Associations between autoimmune conditions and hepatobiliary cancer risk among elderly US adults. *International Journal of Cancer*, **2019**, 144, 707-717	7.5	10
503	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	6.2	9
502	Second primary cancer after female breast cancer: Familial risks and cause of death. *Cancer Medicine*, **2019**, 8, 400-407	4.8	9
501	Genetic variation associated with chromosomal aberration frequency: A genome-wide association study. *Environmental and Molecular Mutagenesis*, **2019**, 60, 17-28	3.2	8
500	Familial Risks Between Urolithiasis and Cancer. *Scientific Reports*, **2018**, 8, 3083	4.9	1
499	Cytogenetic aberrations in multiple myeloma are associated with shifts in serum immunoglobulin isotypes distribution and levels. *Haematologica*, **2018**, 103, e162-e164	6.6	3

498	RE: Familial Cancer Clustering of Urothelial Cancer: A Population-Based Case-Control Study. *Journal of the National Cancer Institute*, **2018**, 110, 1277-1278	9.7	1
497	Genome-wide association study identifies susceptibility loci for B-cell childhood acute lymphoblastic leukemia. *Nature Communications*, **2018**, 9, 1340	17.4	39
496	The multiple myeloma risk allele at 5q15 lowers ELL2 expression and increases ribosomal gene expression. *Nature Communications*, **2018**, 9, 1649	17.4	15
495	Whole genome sequencing reveals DICER1 as a candidate predisposing gene in familial Hodgkin lymphoma. *International Journal of Cancer*, **2018**, 143, 2076-2078	7.5	9
494	Bortezomib-induced peripheral neuropathy: A genome-wide association study on multiple myeloma patients. *Hematological Oncology*, **2018**, 36, 232-237	1.3	14
493	Bleomycin-induced chromosomal damage and shortening of telomeres in peripheral blood lymphocytes of incident cancer patients. *Genes Chromosomes and Cancer*, **2018**, 57, 61-69	5	10
492	Familial Urinary Bladder Cancer with Other Cancers. *European Urology Oncology*, **2018**, 1, 461-466	6.7	4
491	Coding variants in NOD-like receptors: An association study on risk and survival of colorectal cancer. *PLoS ONE*, **2018**, 13, e0199350	3.7	5
490	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	14.6	10
489	Familial risks in and between stone diseases: sialolithiasis, urolithiasis and cholelithiasis in the population of Sweden. *BMC Nephrology*, **2018**, 19, 158	2.7	2
488	Familial risks of acute myeloid leukemia, myelodysplastic syndromes, and myeloproliferative neoplasms. *Blood*, **2018**, 132, 973-976	2.2	19
487	Borderline Ovarian Tumors Share Familial Risks with Themselves and Invasive Cancers. *Cancer Epidemiology Biomarkers and Prevention*, **2018**, 27, 1358-1363	4	0
486	Familial Associations in Testicular Cancer with Other Cancers. *Scientific Reports*, **2018**, 8, 10880	4.9	9
485	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	3	5
484	HLA and KIR Associations of Cervical Neoplasia. *Journal of Infectious Diseases*, **2018**, 218, 2006-2015	7	15
483	Chemotherapy-induced peripheral neuropathy: evidence from genome-wide association studies and replication within multiple myeloma patients. *BMC Cancer*, **2018**, 18, 820	4.8	11
482	Familial Ovarian Cancer Clusters with Other Cancers. *Scientific Reports*, **2018**, 8, 11561	4.9	4
481	Multiple myeloma: family history and mortality in second primary cancers. *Blood Cancer Journal*, **2018**, 8, 75	7	5

480	Familial Cancer Variant Prioritization Pipeline version 2 (FCVPPv2) applied to a papillary thyroid cancer family. *Scientific Reports*, **2018**, 8, 11635	4.9	15
479	Investigation of single and synergic effects of NLRC5 and PD-L1 variants on the risk of colorectal cancer. *PLoS ONE*, **2018**, 13, e0192385	3.7	16
478	Single nucleotide polymorphisms within Mucin-type O-glycan genes are associated with colorectal cancer survival.. *Journal of Clinical Oncology*, **2018**, 36, e15607-e15607	2.2	
477	Genetic correlation between multiple myeloma and chronic lymphocytic leukaemia provides evidence for shared aetiology. *Blood Cancer Journal*, **2018**, 9, 1	7	18
476	Familial risks in urolithiasis in the population of Sweden. *BJU International*, **2018**, 121, 479-485	5.6	11
475	Response: Methods for second primary cancers evaluation have to be standardized. *International Journal of Cancer*, **2018**, 142, 1286-1287	7.5	
474	Familial Risks and Mortality in Second Primary Cancers in Melanoma. *JNCI Cancer Spectrum*, **2018**, 2, pky068	4.8	8
473	Familial risks of ovarian cancer by age at diagnosis, proband type and histology. *PLoS ONE*, **2018**, 13, e0205000	3.7	9
472	Importance of tumor location and histology in familial risk of upper gastrointestinal cancers: a nationwide cohort study. *Clinical Epidemiology*, **2018**, 10, 1169-1179	5.9	9
471	Prostate cancer survivors: Risk and mortality in second primary cancers. *Cancer Medicine*, **2018**, 7, 5752-5759	4.8	11
470	Second primary cancers in non-Hodgkin lymphoma: Bidirectional analyses suggesting role for immune dysfunction. *International Journal of Cancer*, **2018**, 143, 2449-2457	7.5	14
469	Familial risks of second primary cancers and mortality in ovarian cancer patients. *Clinical Epidemiology*, **2018**, 10, 1457-1466	5.9	6
468	Clinical landscape of cancer metastases. *Cancer Medicine*, **2018**, 7, 5534-5542	4.8	27
467	Identification of multiple risk loci and regulatory mechanisms influencing susceptibility to multiple myeloma. *Nature Communications*, **2018**, 9, 3707	17.4	57
466	Genome-wide association study implicates immune dysfunction in the development of Hodgkin lymphoma. *Blood*, **2018**, 132, 2040-2052	2.2	10
465	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	6.2	6
464	Familial risk of pleural mesothelioma increased drastically in certain occupations: A nationwide prospective cohort study. *European Journal of Cancer*, **2018**, 103, 1-6	7.5	7
463	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	2.2	3

462	Genetic variation of acquired structural chromosomal aberrations. *Mutation Research - Genetic Toxicology and Environmental Mutagenesis*, **2018**, 836, 13-21	3	15
461	Genome-wide association analysis of chronic lymphocytic leukaemia, Hodgkin lymphoma and multiple myeloma identifies pleiotropic risk loci. *Scientific Reports*, **2017**, 7, 41071	4.9	27
460	Risk of other Cancers in Families with Melanoma: Novel Familial Links. *Scientific Reports*, **2017**, 7, 42601	4.9	17
459	Common cancers share familial susceptibility: implications for cancer genetics and counselling. *Journal of Medical Genetics*, **2017**, 54, 248-253	5.8	8
458	Functional germline variants in driver genes of breast cancer. *Cancer Causes and Control*, **2017**, 28, 259-27[illegible]	2.8	7
457	Low expression of hexokinase-2 is associated with false-negative FDG-positron emission tomography in multiple myeloma. *Blood*, **2017**, 130, 30-34	2.2	120
456	Genetics of gallbladder cancer. *Lancet Oncology, The*, **2017**, 18, e296	21.7	6
455	Genomewide association study on monoclonal gammopathy of unknown significance (MGUS). *European Journal of Haematology*, **2017**, 99, 70-79	3.8	13
454	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	6.6	14
453	Concordant and discordant familial cancer: Familial risks, proportions and population impact. *International Journal of Cancer*, **2017**, 140, 1510-1516	7.5	40
452	Risk of Second Cancer in Hodgkin Lymphoma Survivors and Influence of Family History. *Journal of Clinical Oncology*, **2017**, 35, 1584-1590	2.2	46
451	Familial associations of male breast cancer with other cancers. *Breast Cancer Research and Treatment*, **2017**, 166, 897-902	4.4	5
450	Genetic Predisposition to Multiple Myeloma at 5q15 Is Mediated by an ELL2 Enhancer Polymorphism. *Cell Reports*, **2017**, 20, 2556-2564	10.6	15
449	Surveillance Bias in Cancer Risk After Unrelated Medical Conditions: Example Urolithiasis. *Scientific Reports*, **2017**, 7, 8073	4.9	13
448	Other cancers in lung cancer families are overwhelmingly smoking-related cancers. *ERJ Open Research*, **2017**, 3,	3.5	3
447	Risk of second primary cancers in women diagnosed with endometrial cancer in German and Swedish cancer registries. *International Journal of Cancer*, **2017**, 141, 2270-2280	7.5	8
446	Familial associations of female breast cancer with other cancers. *International Journal of Cancer*, **2017**, 141, 2253-2259	7.5	13
445	Genome-wide association study of classical Hodgkin lymphoma identifies key regulators of disease susceptibility. *Nature Communications*, **2017**, 8, 1892	17.4	24

444	Familial Associations of Colorectal Cancer with Other Cancers. *Scientific Reports*, **2017**, 7, 5243	4.9	5
443	Genome-wide association study of clinical parameters in immunoglobulin light chain amyloidosis in three patient cohorts. *Haematologica*, **2017**, 102, e411-e414	6.6	7
442	Whole-exome sequencing identifies novel candidate predisposition genes for familial polycythemia vera. *Human Genomics*, **2017**, 11, 6	6.8	9
441	Familial Associations Between Prostate Cancer and Other Cancers. *European Urology*, **2017**, 71, 162-165	10.2	13
440	Identification of miRSNPs associated with the risk of multiple myeloma. *International Journal of Cancer*, **2017**, 140, 526-534	7.5	6
439	Genetic Susceptibility to Bortezomib-Induced Peripheral Neuroropathy: Replication of the Reported Candidate Susceptibility Loci. *Neurochemical Research*, **2017**, 42, 925-931	4.6	12
438	Familial risks for gallstones in the population of Sweden. *BMJ Open Gastroenterology*, **2017**, 4, e000188	3.9	6
437	Direct evidence for a polygenic etiology in familial multiple myeloma. *Blood Advances*, **2017**, 1, 619-623	7.8	13
436	Defining the genetic susceptibility to cervical neoplasia-A genome-wide association study. *PLoS Genetics*, **2017**, 13, e1006866	6	55
435	Inherited variants in genes somatically mutated in thyroid cancer. *PLoS ONE*, **2017**, 12, e0174995	3.7	4
434	Impact of functional germline variants and a deletion polymorphism in APOBEC3A and APOBEC3B on breast cancer risk and survival in a Swedish study population. *Journal of Cancer Research and Clinical Oncology*, **2016**, 142, 273-6	4.9	21
433	Epidemiology, Risk Factors, and Survival in CUP: Pointers to Disease Mechanisms **2016**, 5-25		
432	The epidemiology of metastases in neuroendocrine tumors. *International Journal of Cancer*, **2016**, 139, 2679-2686	7.5	132
431	Multiple myeloma risk variant at 7p15.3 creates an IRF4-binding site and interferes with CDCA7L expression. *Nature Communications*, **2016**, 7, 13656	17.4	26
430	Genome-wide association study identifies multiple susceptibility loci for multiple myeloma. *Nature Communications*, **2016**, 7, 12050	17.4	101
429	Pedigree based DNA sequencing pipeline for germline genomes of cancer families. *Hereditary Cancer in Clinical Practice*, **2016**, 14, 16	2.3	5
428	Patterns of metastasis in colon and rectal cancer. *Scientific Reports*, **2016**, 6, 29765	4.9	409
427	Risk of Second Primary Cancers in Multiple Myeloma Survivors in German and Swedish Cancer Registries. *Scientific Reports*, **2016**, 6, 22084	4.9	12

426	Age-Dependent Metastatic Spread and Survival: Cancer of Unknown Primary as a Model. *Scientific Reports*, **2016**, 6, 23725	4.9	9
425	Risk of second primary cancers after malignant mesothelioma and vice versa. *Cancer Letters*, **2016**, 379, 94-9	9.9	7
424	Survival in familial and non-familial breast cancer by age and stage at diagnosis. *European Journal of Cancer*, **2016**, 52, 10-8	7.5	7
423	A Comprehensive Meta-analysis of Case-Control Association Studies to Evaluate Polymorphisms Associated with the Risk of Differentiated Thyroid Carcinoma. *Cancer Epidemiology Biomarkers and Prevention*, **2016**, 25, 700-13	4	16
422	Evidence of Inbreeding in Hodgkin Lymphoma. *PLoS ONE*, **2016**, 11, e0154259	3.7	7
421	A common variant within the HNF1B gene is associated with overall survival of multiple myeloma patients: results from the IMMEnSE consortium and meta-analysis. *Oncotarget*, **2016**, 7, 59029-59048	3.3	14
420	Mapping of deletion breakpoints at the CDKN2A locus in melanoma: detection of MTAP-ANRIL fusion transcripts. *Oncotarget*, **2016**, 7, 16490-504	3.3	18
419	Germline genetics of cancer of unknown primary (CUP) and its specific subtypes. *Oncotarget*, **2016**, 7, 22140-9	3.3	7
418	Metastatic spread in patients with gastric cancer. *Oncotarget*, **2016**, 7, 52307-52316	3.3	158
417	Origin of B-Cell Neoplasms in Autoimmune Disease. *PLoS ONE*, **2016**, 11, e0158360	3.7	13
416	TERT promoter mutations in melanoma survival. *International Journal of Cancer*, **2016**, 139, 75-84	7.5	79
415	Cancer of unknown primary is associated with diabetes. *European Journal of Cancer Prevention*, **2016**, 25, 246-51	2	9
414	Location of metastases in cancer of unknown primary are not random and signal familial clustering. *Scientific Reports*, **2016**, 6, 22891	4.9	13
413	The Incidence of Senile Cataract and Glaucoma is Increased in Patients with Plasma Cell Dyscrasias: Etiologic Implications. *Scientific Reports*, **2016**, 6, 28500	4.9	9
412	Predictive and Prognostic Clinical Variables in Cancer Patients Treated With Adenoviral Oncolytic Immunotherapy. *Molecular Therapy*, **2016**, 24, 1323-32	11.7	19
411	Analysis of functional germline variants in APOBEC3 and driver genes on breast cancer risk in Moroccan study population. *BMC Cancer*, **2016**, 16, 165	4.8	16
410	Polymorphisms within base and nucleotide excision repair pathways and risk of differentiated thyroid carcinoma. *DNA Repair*, **2016**, 41, 27-31	4.3	5
409	Runs of homozygosity and inbreeding in thyroid cancer. *BMC Cancer*, **2016**, 16, 227	4.8	15

408	Genetic variation in the major mitotic checkpoint genes associated with chromosomal aberrations in healthy humans. *Cancer Letters*, **2016**, 380, 442-446	9.9	8
407	Risk of Next Melanoma in Patients With Familial and Sporadic Melanoma by Number of Previous Melanomas. *JAMA Dermatology*, **2015**, 151, 607-15	5.1	21
406	Heritability estimates on Hodgkin's lymphoma: a genomic- versus population-based approach. *European Journal of Human Genetics*, **2015**, 23, 824-30	5.3	9
405	Case-control estimation of the impact of oncolytic adenovirus on the survival of patients with refractory solid tumors. *Molecular Therapy*, **2015**, 23, 321-9	11.7	10
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403	A simple-to-use method incorporating genomic markers into prostate cancer risk prediction tools facilitated future validation. *Journal of Clinical Epidemiology*, **2015**, 68, 563-73	5.7	6
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