D Timothy Bishop

List of Publications by Year
in descending order

304	32,854	4653 85	4427 172
papers	citations	h-index	g-index

319	319	319	26921
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Identification of the breast cancer susceptibility gene BRCA2. Nature, 1995, 378, 789-792.	13.7	3,230
2	Genetic Heterogeneity and Penetrance Analysis of the BRCA1 and BRCA2 Genes in Breast Cancer Families. American Journal of Human Genetics, 1998, 62, 676-689.	2.6	2,662
3	Risks of cancer in BRCA1-mutation carriers. Lancet, The, 1994, 343, 692-695.	6.3	1,764
4	Long-term effect of aspirin on cancer risk in carriers of hereditary colorectal cancer: an analysis from the CAPP2 randomised controlled trial. Lancet, The, 2011, 378, 2081-2087.	6.3	849
5	Gene for von Recklinghausen neurofibromatosis is in the pericentromeric region of chromosome 17. Science, 1987, 236, 1100-1102.	6.0	687
6	Multifactorial Analysis of Differences Between Sporadic Breast Cancers and Cancers Involving BRCA1 and BRCA2 Mutations. Journal of the National Cancer Institute, 1998, 90, 1138-1145.	3.0	652
7	Prediction of BRCA1 Status in Patients with Breast Cancer Using Estrogen Receptor and Basal Phenotype. Clinical Cancer Research, 2005, 11, 5175-5180.	3.2	577
8	Sequence variants at the TERT-CLPTM1L locus associate with many cancer types. Nature Genetics, 2009, 41, 221-227.	9.4	572
9	A genome-wide association study identifies colorectal cancer susceptibility loci on chromosomes 10p14 and 8q23.3. Nature Genetics, 2008, 40, 623-630.	9.4	514
10	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. Nature Genetics, 2010, 42, 978-984.	9.4	493
11	Common Inheritance of Susceptibility to Colonic Adenomatous Polyps and Associated Colorectal Cancers. New England Journal of Medicine, 1988, 319, 533-537.	13.9	464
12	Geographical Variation in the Penetrance of CDKN2A Mutations for Melanoma. Journal of the National Cancer Institute, 2002, 94, 894-903.	3.0	435
13	Genome-wide association study identifies three loci associated with melanoma risk. Nature Genetics, 2009, 41, 920-925.	9.4	422
14	A novel recurrent mutation in MITF predisposes to familial and sporadic melanoma. Nature, 2011, 480, 99-103.	13.7	413
15	Sequence variant on 8q24 confers susceptibility to urinary bladder cancer. Nature Genetics, 2008, 40, 1307-1312.	9.4	377
16	Discovery of common and rare genetic risk variants for colorectal cancer. Nature Genetics, 2019, 51, 76-87.	9.4	377
17	High-risk Melanoma Susceptibility Genes and Pancreatic Cancer, Neural System Tumors, and Uveal Melanoma across GenoMEL. Cancer Research, 2006, 66, 9818-9828.	0.4	373
18	Features associated with germline CDKN2A mutations: a GenoMEL study of melanoma-prone families from three continents. Journal of Medical Genetics, 2006, 44, 99-106.	1.5	350

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19	Construction of linkage maps with DNA markers for human chromosomes. Nature, 1985, 313, 101-105.	13.7	336
20	Genetic variation in the prostate stem cell antigen gene PSCA confers susceptibility to urinary bladder cancer. Nature Genetics, 2009, 41, 991-995.	9.4	321
21	Risk of Colorectal Cancer in the Families of Patients with Adenomatous Polyps. New England Journal of Medicine, 1996, 334, 82-87.	13.9	320
22	Variants near DMRT1, TERT and ATF7IP are associated with testicular germ cell cancer. Nature Genetics, 2010, 42, 604-607.	9.4	320
23	POT1 loss-of-function variants predispose to familial melanoma. Nature Genetics, 2014, 46, 478-481.	9.4	319
24	A genome-wide association study of testicular germ cell tumor. Nature Genetics, 2009, 41, 807-810.	9.4	317
25	Structure of the Human MSH2 Locus and Analysis of Two Muir-Torre Kindreds for msh2 Mutations. Genomics, 1994, 24, 516-526.	1.3	276
26	Effect of Aspirin or Resistant Starch on Colorectal Neoplasia in the Lynch Syndrome. New England Journal of Medicine, 2008, 359, 2567-2578.	13.9	273
27	Serum 25-Hydroxyvitamin D₃ Levels Are Associated With Breslow Thickness at Presentation and Survival From Melanoma. Journal of Clinical Oncology, 2009, 27, 5439-5444.	0.8	263
28	Localization to Xq27 of a susceptibility gene for testicular germ-cell tumours. Nature Genetics, 2000, 24, 197-200.	9.4	260
29	New loci for body fat percentage reveal link between adiposity and cardiometabolic disease risk. Nature Communications, 2016, 7, 10495.	5.8	245
30	A pharmacogenetic study to investigate the role of dietary carcinogens in the etiology of colorectal cancer. Carcinogenesis, 2002, 23, 1839-1850.	1.3	241
31	Dominant Inheritance of Adenomatous Colonic Polyps and Colorectal Cancer. New England Journal of Medicine, 1985, 312, 1540-1544.	13.9	240
32	Genome-wide association study identifies three new melanoma susceptibility loci. Nature Genetics, 2011, 43, 1108-1113.	9.4	230
33	Pathology of Ovarian Cancers in BRCA1 and BRCA2 Carriers. Clinical Cancer Research, 2004, 10, 2473-2481.	3.2	224
34	Cancer prevention with aspirin in hereditary colorectal cancer (Lynch syndrome), 10-year follow-up and registry-based 20-year data in the CAPP2 study: a double-blind, randomised, placebo-controlled trial. Lancet, The, 2020, 395, 1855-1863.	6.3	220
35	Sun exposure and melanoma risk at different latitudes: a pooled analysis of 5700 cases and 7216 controls. International Journal of Epidemiology, 2009, 38, 814-830.	0.9	219
36	Genome-wide meta-analysis identifies five new susceptibility loci for cutaneous malignant melanoma. Nature Genetics, 2015, 47, 987-995.	9.4	218

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37	Genome-wide association study identifies variants at 9p21 and 22q13 associated with development of cutaneous nevi. Nature Genetics, 2009, 41, 915-919.	9.4	204
38	The Y Deletion gr/gr and Susceptibility to Testicular Germ Cell Tumor. American Journal of Human Genetics, 2005, 77, 1034-1043.	2.6	197
39	A Randomized Placebo-Controlled Prevention Trial of Aspirin and/or Resistant Starch in Young People with Familial Adenomatous Polyposis. Cancer Prevention Research, 2011, 4, 655-665.	0.7	193
40	Physical activity and risks of breast and colorectal cancer: a Mendelian randomisation analysis. Nature Communications, 2020, 11, 597.	5.8	193
41	MRE11 Expression Is Predictive of Cause-Specific Survival following Radical Radiotherapy for Muscle-Invasive Bladder Cancer. Cancer Research, 2010, 70, 7017-7026.	0.4	184
42	Hereditary Hemochromatosis: Analysis of Laboratory Expression of the Disease by Genotype in 18 Pedigrees. American Journal of Clinical Pathology, 1982, 78, 196-207.	0.4	181
43	A sequence variant at 4p16.3 confers susceptibility to urinary bladder cancer. Nature Genetics, 2010, 42, 415-419.	9.4	169
44	RHBDF2 Mutations Are Associated with Tylosis, a Familial Esophageal Cancer Syndrome. American Journal of Human Genetics, 2012, 90, 340-346.	2.6	162
45	Increased Risk of Cancer Other Than Melanoma in CDKN2A Founder Mutation (p16-Leiden)-Positive Melanoma Families. Clinical Cancer Research, 2008, 14, 7151-7157.	3.2	161
46	Meta-analysis identifies four new loci associated with testicular germ cell tumor. Nature Genetics, 2013, 45, 680-685.	9.4	154
47	The information content of phase-known matings for ordering genetic loci. Genetic Epidemiology, 1985, 2, 349-361.	0.6	153
48	Haplotype and Phenotype Analysis of Nine Recurrent BRCA2 Mutations in 111 Families: Results of an International Study. American Journal of Human Genetics, 1998, 62, 1381-1388.	2.6	150
49	Identification of nine new susceptibility loci for testicular cancer, including variants near DAZL and PRDM14. Nature Genetics, 2013, 45, 686-689.	9.4	149
50	Population-based family studies in genetic epidemiology. Lancet, The, 2005, 366, 1397-1406.	6.3	148
51	Genome-wide association study identifies a new melanoma susceptibility locus at 1q21.3. Nature Genetics, 2011, 43, 1114-1118.	9.4	140
52	Genome-wide linkage screen for testicular germ cell tumour susceptibility loci. Human Molecular Genetics, 2006, 15, 443-451.	1.4	138
53	Genome-wide association meta-analyses combining multiple risk phenotypes provide insights into the genetic architecture of cutaneous melanoma susceptibility. Nature Genetics, 2020, 52, 494-504.	9.4	138
54	Germline Mutations of the CDKN2 Gene in UK Melanoma Families. Human Molecular Genetics, 1997, 6, 2061-2067.	1.4	135

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55	European genome-wide association study identifies SLC14A1 as a new urinary bladder cancer susceptibility gene. Human Molecular Genetics, 2011, 20, 4268-4281.	1.4	134
56	Nonsense Mutations in the Shelterin Complex Genes ACD and TERF2IP in Familial Melanoma. Journal of the National Cancer Institute, 2015, 107, .	3.0	134
57	Vitamin D receptor gene polymorphisms, serum 25-hydroxyvitamin D levels, and melanoma: UK caseâ€“control comparisons and a meta-analysis of published VDR data. European Journal of Cancer, 2009, 45, 3271-3281.	1.3	127
58	People of the British Isles: preliminary analysis of genotypes and surnames in a UK-control population. European Journal of Human Genetics, 2012, 20, 203-210.	1.4	126
59	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. American Journal of Human Genetics, 2020, 107, 432-444.	2.6	124
60	Rare disruptive mutations and their contribution to the heritable risk of colorectal cancer. Nature Communications, 2016, 7, 11883.	5.8	122
61	Identification of 19 new risk loci and potential regulatory mechanisms influencing susceptibility to testicular germ cell tumor. Nature Genetics, 2017, 49, 1133-1140.	9.4	120
62	Quantitating Genetic and Nongenetic Factors Influencing Androgen Production and Clearance Rates in Men*. Journal of Clinical Endocrinology and Metabolism, 1988, 67, 104-109.	1.8	116
63	Cardiometabolic effects of genetic upregulation of the interleukin 1 receptor antagonist: a Mendelian randomisation analysis. Lancet Diabetes and Endocrinology,the, 2015, 3, 243-253.	5.5	115
64	IRF4 Variants Have Age-Specific Effects on Nevus Count and Predispose to Melanoma. American Journal of Human Genetics, 2010, 87, 6-16.	2.6	114
65	Localization of a Novel Melanoma Susceptibility Locus to 1p22. American Journal of Human Genetics, 2003, 73, 301-313.	2.6	113
66	The effect of nutritional factors on sex hormone levels in male twins. Genetic Epidemiology, 1988, 5, 43-59.	0.6	111
67	A variant in FTO shows association with melanoma risk not due to BMI. Nature Genetics, 2013, 45, 428-432.	9.4	111
68	Cumulative Burden of Colorectal Cancerâ€“Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. Gastroenterology, 2020, 158, 1274-1286.e12.	0.6	110
69	The Effect on Melanoma Risk of Genes Previously Associated With Telomere Length. Journal of the National Cancer Institute, 2014, 106, .	3.0	109
70	Polymorphisms in DNA Repair Genes, Smoking, and Bladder Cancer Risk: Findings from the International Consortium of Bladder Cancer. Cancer Research, 2009, 69, 6857-6864.	0.4	107
71	A Genomewide Linkage Study of 1,933 Families Affected by Premature Coronary Artery Disease: The British Heart Foundation (BHF) Family Heart Study. American Journal of Human Genetics, 2005, 77, 1011-1020.	2.6	105
72	Meta-analysis of five genome-wide association studies identifies multiple new loci associated with testicular germ cell tumor. Nature Genetics, 2017, 49, 1141-1147.	9.4	105

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73	The role of microRNA-binding site polymorphisms in DNA repair genes as risk factors for bladder cancer and breast cancer and their impact on radiotherapy outcomes. Carcinogenesis, 2012, 33, 581-586.	1.3	103
74	Germline MC1R status influences somatic mutation burden in melanoma. Nature Communications, 2016, 7, 12064.	5.8	103
75	Genotype/Phenotype and Penetrance Studies in Melanoma Families with Germline CDKN2A Mutations. Journal of Investigative Dermatology, 2000, 114, 28-33.	0.3	102
76	Melanocytic Nevi, Nevus Genes, and Melanoma Risk in a Large Case-Control Study in the United Kingdom. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 2043-2054.	1.1	102
77	Genome-wide association study identifies a common variant in RAD51B associated with male breast cancer risk. Nature Genetics, 2012, 44, 1182-1184.	9.4	99
78	Long-term effect of resistant starch on cancer risk in carriers of hereditary colorectal cancer: an analysis from the CAPP2 randomised controlled trial. Lancet Oncology, The, 2012, 13, 1242-1249.	5.1	95
79	Frequent p16-Independent Inactivation of p14ARF in Human Melanoma. Journal of the National Cancer Institute, 2008, 100, 784-795.	3.0	94
80	Gene Expression Profiling of Paraffin-Embedded Primary Melanoma Using the DASL Assay Identifies Increased Osteopontin Expression as Predictive of Reduced Relapse-Free Survival. Clinical Cancer Research, 2009, 15, 6939-6946.	3.2	93
81	The introduction of SCF3 into aromatic substrates using CuSCF3 and alumina-supported CuSCF3. Journal of Fluorine Chemistry, 1990, 48, 249-253.	0.9	91
82	Meta- and Pooled Analyses of the Methylenetetrahydrofolate Reductase (MTHFR) C677T Polymorphism and Colorectal Cancer: A HuGE-GSEC Review. American Journal of Epidemiology, 2009, 170, 1207-1221.	1.6	91
83	Obesity, Aspirin, and Risk of Colorectal Cancer in Carriers of Hereditary Colorectal Cancer: A Prospective Investigation in the CAPP2 Study. Journal of Clinical Oncology, 2015, 33, 3591-3597.	0.8	91
84	Relationship between sun exposure and melanoma risk for tumours in different body sites in a large case-control study in a temperate climate. European Journal of Cancer, 2011, 47, 732-741.	1.3	90
85	Circulating Levels of Insulin-like Growth Factor 1 and Insulin-like Growth Factor Binding Protein 3 Associate With Risk of Colorectal Cancer Based on Serologic and Mendelian Randomization Analyses. Gastroenterology, 2020, 158, 1300-1312.e20.	0.6	90
86	Measures of familial aggregation depend on definition of family history: meta-analysis for colorectal cancer. Journal of Clinical Epidemiology, 2006, 59, 114-124.	2.4	89
87	How common is the atypical mole syndrome phenotype in apparently sporadic melanoma?. Journal of the American Academy of Dermatology, 1993, 29, 989-996.	0.6	88
88	Investigation of interaction between N-acetyltransferase 2 and heterocyclic amines as potential risk factors for colorectal cancer. Carcinogenesis, 2003, 24, 275-282.	1.3	88
89	Novel pleiotropic risk loci for melanoma and nevus density implicate multiple biological pathways. Nature Communications, 2018, 9, 4774.	5.8	87
90	A pooled analysis of melanocytic nevus phenotype and the risk of cutaneous melanoma at different latitudes. International Journal of Cancer, 2009, 124, 420-428.	2.3	84

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91	Somatic KIT mutations occur predominantly in seminoma germ cell tumors and are not predictive of bilateral disease: Report of 220 tumors and review of literature. Genes Chromosomes and Cancer, 2008, 47, 34-42.	1.5	83
92	Adiposity, metabolites, and colorectal cancer risk: Mendelian randomization study. BMC Medicine, 2020, 18, 396.	2.3	76
93	Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. Nature Communications, 2020, 11, 3353.	5.8	75
94	Genetic analysis of the BRCA1 region in a large breast/ovarian family: refinement of the minimal region containing BRCA1. Human Molecular Genetics, 1993, 2, 1823-1828.	1.4	71
95	Tylosis oesophageal cancer mapped. Nature Genetics, 1994, 8, 319-321.	9.4	71
96	Meta Association of Colorectal Cancer Confirms Risk Alleles at 8q24 and 18q21. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 616-621.	1.1	71
97	Î²-Cateninâ€“mediated immune evasion pathway frequently operates in primary cutaneous melanomas. Journal of Clinical Investigation, 2018, 128, 2048-2063.	3.9	71
98	A mutation hotspot at the p14ARF splice site. Oncogene, 2005, 24, 4604-4608.	2.6	70
99	The Genetic Epidemiology of Early-Onset Epithelial Ovarian Cancer: A Population-Based Study. American Journal of Human Genetics, 1999, 65, 1725-1732.	2.6	69
100	Cell-typeâ€“specific eQTL of primary melanocytes facilitates identification of melanoma susceptibility genes. Genome Research, 2018, 28, 1621-1635.	2.4	67
101	Vegetable, fruit and meat consumption and potential risk modifying genes in relation to colorectal cancer. International Journal of Cancer, 2004, 112, 259-264.	2.3	65
102	Vitamin Dâ€“VDR Signaling Inhibits Wnt/Î²-Cateninâ€“Mediated Melanoma Progression and Promotes Antitumor Immunity. Cancer Research, 2019, 79, 5986-5998.	0.4	65
103	Family studies in melanoma: identification of the atypical mole syndrome (AMS) phenotype. Melanoma Research, 1994, 4, 199-206.	0.6	62
104	The Effect of Sun Exposure in Determining Nevus Density in UK Adolescent Twins. Journal of Investigative Dermatology, 2005, 124, 56-62.	0.3	62
105	Refinement of the basis and impact of common 11q23.1 variation to the risk of developing colorectal cancer. Human Molecular Genetics, 2008, 17, 3720-3727.	1.4	61
106	Inherited variants in the MC1R gene and survival from cutaneous melanoma: a BioGenoMEL study. Pigment Cell and Melanoma Research, 2012, 25, 384-394.	1.5	61
107	25â€Hydroxyvitamin D₂/D₃ levels and factors associated with systemic inflammation and melanoma survival in the Leeds Melanoma Cohort. International Journal of Cancer, 2015, 136, 2890-2899.	2.3	61
108	Management of familial melanoma. Lancet Oncology, The, 2007, 8, 46-54.	5.1	60

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109	Mutation screening of theCDKN2A promoter in melanoma families. , 2000, 28, 45-57.		59
110	Testicular germ cell tumor susceptibility associated with the UCK2 locus on chromosome 1q23. Human Molecular Genetics, 2013, 22, 2748-2753.	1.4	59
111	Mutation testing in melanoma families: INK4A, CDK4 and INK4D. British Journal of Cancer, 1999, 80, 295-300.	2.9	57
112	Chemoprevention in Lynch syndrome. Familial Cancer, 2013, 12, 707-718.	0.9	57
113	AT-tributable risks?. Nature Genetics, 1997, 15, 226-226.	9.4	56
114	Melanoma sentinel node biopsy and prediction models for relapse and overall survival. British Journal of Cancer, 2010, 103, 1229-1236.	2.9	54
115	Large-scale Sequencing of Testicular Germ Cell Tumour (TGCT) Cases Excludes Major TGCT Predisposition Gene. European Urology, 2018, 73, 828-831.	0.9	54
116	Patterns of Expression of DNA Repair Genes and Relapse From Melanoma. Clinical Cancer Research, 2010, 16, 5211-5221.	3.2	53
117	Development and External Validation of a Melanoma Risk Prediction Model Based on Self-assessed Risk Factors. JAMA Dermatology, 2016, 152, 889.	2.0	53
118	Evidence for a colorectal cancer susceptibility locus on chromosome 3q21-q24 from a high-density SNP genome-wide linkage scan. Human Molecular Genetics, 2006, 15, 2903-2910.	1.4	52
119	Assessing the Incremental Contribution of Common Genomic Variants to Melanoma Risk Prediction in Two Population-Based Studies. Journal of Investigative Dermatology, 2018, 138, 2617-2624.	0.3	52
120	Localisation of susceptibility genes for familial testicular germ cell tumour. Apmis, 2003, 111, 128-135.	0.9	51
121	Evidence of Linkage to Chromosome 9q22.33 in Colorectal Cancer Kindreds from the United Kingdom. Cancer Research, 2006, 66, 5003-5006.	0.4	51
122	Testicular microlithiasis as a familial risk factor for testicular germ cell tumour. British Journal of Cancer, 2007, 97, 1701-1706.	2.9	51
123	Germline TERT promoter mutations are rare in familial melanoma. Familial Cancer, 2016, 15, 139-144.	0.9	51
124	A common intronic variant of PARP1 confers melanoma risk and mediates melanocyte growth via regulation of MITF. Nature Genetics, 2017, 49, 1326-1335.	9.4	51
125	Identification of a modifier gene locus on chromosome 1p35-36 in familial adenomatous polyposis. Human Genetics, 1997, 99, 653-657.	1.8	49
126	Association between hormonal genetic polymorphisms and early-onset prostate cancer. Prostate Cancer and Prostatic Diseases, 2005, 8, 95-102.	2.0	49

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127	Constraints on the Progenitor of SN 2016gkg from Its Shock-cooling Light Curve. Astrophysical Journal Letters, 2017, 837, L2.	3.0	49
128	Candidate regions for testicular cancer susceptibility genes. Apmis, 1998, 106, 64-72.	0.9	48
129	Close mapping of the focal non-epidermolytic palmoplantar keratoderma (PPK) locus associated with oesophageal cancer (TOC). Human Molecular Genetics, 1996, 5, 857-860.	1.4	47
130	Ki-rasgene mutations are invariably present in low-grade mucinous tumors of the vermiform appendix. Scandinavian Journal of Gastroenterology, 2011, 46, 869-874.	0.6	47
131	Deletion at chromosome arm 9p in relation to BRAF/NRAS mutations and prognostic significance for primary melanoma. Genes Chromosomes and Cancer, 2010, 49, 425-438.	1.5	46
132	Genome-wide association study yields variants at 20p12.2 that associate with urinary bladder cancer. Human Molecular Genetics, 2014, 23, 5545-5557.	1.4	46
133	p53 Protein Detected By Immunohistochemical Staining is Not Always Mutant. Disease Markers, 1993, 11, 239-250.	0.6	45
134	The polyAT, intronic IVS11-6 and Lys939Gln XPC polymorphisms are not associated with transitional cell carcinoma of the bladder. British Journal of Cancer, 2005, 92, 2262-2265.	2.9	45
135	Prevalence and predictors of germline CDKN2A mutations for melanoma cases from Australia, Spain and the United Kingdom. Hereditary Cancer in Clinical Practice, 2014, 12, 20.	0.6	45
136	Genetic and Environmental Determinants of Immune Response to Cutaneous Melanoma. Cancer Research, 2019, 79, 2684-2696.	0.4	45
137	MLH1 â^'93G>A promoter polymorphism and risk of mismatch repair deficient colorectal cancer. International Journal of Cancer, 2008, 123, 2456-2459.	2.3	44
138	Genetic architectures of proximal and distal colorectal cancer are partly distinct. Gut, 2021, 70, 1325-1334.	6.1	44
139	Independent replication of a melanoma subtype gene signature and evaluation of its prognostic value and biological correlates in a population cohort. Oncotarget, 2015, 6, 11683-11693.	0.8	44
140	Mutations and alternative splicing of theBRCA1 gene in UK breast/ovarian cancer families. , 1997, 18, 102-110.		43
141	AfterhMSH2 andhMLH1?what next? Analysis of three-generational, population-based, early-onset colorectal cancer families. International Journal of Cancer, 2002, 102, 166-171.	2.3	43
142	GSTM1 and CYP1A1 Polymorphisms, Tobacco, Air Pollution, and Lung Cancer: A Study in Rural Thailand. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 667-674.	1.1	42
143	Polymorphisms in xenobiotic metabolizing enzymes and diet influence colorectal adenoma risk. Pharmacogenetics and Genomics, 2010, 20, 315-326.	0.7	42
144	The International Testicular Cancer Linkage Consortium: A clinicopathologic descriptive analysis of 461 familial malignant testicular germ cell tumor kindred. Urologic Oncology: Seminars and Original Investigations, 2010, 28, 492-499.	0.8	42

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145	Melanoma risk for CDKN2A mutation carriers who are relatives of population-based case carriers in Australia and the UK. Journal of Medical Genetics, 2011, 48, 266-272.	1.5	41
146	Identification of a melanoma susceptibility locus and somatic mutation inTET2. Carcinogenesis, 2014, 35, 2097-2101.	1.3	41
147	Next-generation sequencing identifies germline MRE11A variants as markers of radiotherapy outcomes in muscle-invasive bladder cancer. Annals of Oncology, 2014, 25, 877-883.	0.6	41
148	Comprehensive Analysis of 22 XPC Polymorphisms and Bladder Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 2537-2541.	1.1	40
149	Overseas Sun Exposure, Nevus Counts, and Premature Skin Aging in Young English Women: A Population-Based Survey. Journal of Investigative Dermatology, 2009, 129, 50-59.	0.3	40
150	Functional characterization of a multi-cancer risk locus on chr5p15.33 reveals regulation of TERT by ZNF148. Nature Communications, 2017, 8, 15034.	5.8	40
151	Lifestyle and blood pressure levels in male twins in Utah. Genetic Epidemiology, 1988, 5, 277-287.	0.6	39
152	Familial risk and genetic susceptibility for breast cancer. Cancer Causes and Control, 1994, 5, 458-470.	0.8	39
153	The Relationship Between the Epidermal Growth Factor (EGF) 5â€²UTR Variant A61G and Melanoma/Nevus Susceptibility. Journal of Investigative Dermatology, 2004, 123, 755-759.	0.3	39
154	Nongenetic Determinants of Risk forÂEarly-Onset Colorectal Cancer. JNCI Cancer Spectrum, 2021, 5, pkab029.	1.4	39
155	Sister chromatid exchange: Variation by age, sex, smoking, and breast cancer status. Cancer Genetics and Cytogenetics, 1983, 9, 289-299.	1.0	38
156	Analysis of variants in DNA damage signalling genes in bladder cancer. BMC Medical Genetics, 2008, 9, 69.	2.1	38
157	Genetic Variants inXRCC2: New Insights Into Colorectal Cancer Tumorigenesis. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 2476-2484.	1.1	38
158	Identification of a novel susceptibility locus at 13q34 and refinement of the 20p12.2 region as a multi-signal locus associated with bladder cancer risk in individuals of European ancestry. Human Molecular Genetics, 2016, 25, 1203-1214.	1.4	38
159	Loss-of-Function Mutations in the Cell-Cycle Control Gene CDKN2A Impact on Glucose Homeostasis in Humans. Diabetes, 2016, 65, 527-533.	0.3	38
160	Characteristics of familial colon cancer in a large population data base. Cancer, 1989, 64, 1971-1975.	2.0	37
161	Linkage Studies of Non-Syndromic Recessive Deafness (NSRD) in a Family Originating from the Mirpur Region of Pakistan Maps DFNB1 Centromeric to D13S175. Human Molecular Genetics, 1996, 5, 169-173.	1.4	37
162	Analysis of the DND1 gene in men with sporadic and familial testicular germ cell tumors. Genes Chromosomes and Cancer, 2008, 47, 247-252.	1.5	37

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163	Genetic linkage studies in non-epidermolytic palmoplantar keratoderma: evidence for heterogeneity. Human Molecular Genetics, 1995, 4, 1021-1025.	1.4	36
164	An Assessment of the CDKN2A Variant Ala148Thr as a Nevus/Melanoma Susceptibility Allele. Journal of Investigative Dermatology, 2002, 119, 961-965.	0.3	36
165	Prevalence of 9p21 deletions in UK melanoma families. Genes Chromosomes and Cancer, 2005, 44, 292-300.	1.5	36
166	Identification of four new susceptibility loci for testicular germ cell tumour. Nature Communications, 2015, 6, 8690.	5.8	36
167	Identifying Novel Susceptibility Genes for Colorectal Cancer Risk From a Transcriptome-Wide Association Study of 125,478 Subjects. Gastroenterology, 2021, 160, 1164-1178.e6.	0.6	36
168	Can and will a sexual diploid population attain an evolutionary stable strategy?. Journal of Theoretical Biology, 1984, 111, 667-686.	0.8	35
169	Hereditary susceptibility to colorectal cancer. Diseases of the Colon and Rectum, 1996, 39, 739-743.	0.7	34
170	Incidence of DNA replication errors in patients with multiple primary cancers. Diseases of the Colon and Rectum, 1998, 41, 765-769.	0.7	34
171	Association study of asthma and atopy traits and chromosome 5q cytokine cluster markers. Clinical and Experimental Allergy, 1998, 28, 141-150.	1.4	34
172	Envoplakin, a Possible Candidate Gene for Focal NEPPK/Esophageal Cancer (TOC): The Integration of Genetic and Physical Maps of the TOC Region on 17q25. Genomics, 1999, 59, 234-242.	1.3	34
173	CDKN2A and CDK4 variants in Latvian melanoma patients: analysis of a clinic-based population. Melanoma Research, 2007, 17, 185-191.	0.6	34
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