Graham G Giles

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genome-wide association study identifies novel breast cancer susceptibility loci. Nature, 2007, 447, 1087-1093.	13.7	2,165
2	Body-Mass Index and Mortality among 1.46 Million White Adults. New England Journal of Medicine, 2010, 363, 2211-2219.	13.9	1,926
3	REVEL: An Ensemble Method for Predicting the Pathogenicity of Rare Missense Variants. American Journal of Human Genetics, 2016, 99, 877-885.	2.6	1,555
4	Cancer risk in 680 000 people exposed to computed tomography scans in childhood or adolescence: data linkage study of 11 million Australians. BMJ, The, 2013, 346, f2360-f2360.	3.0	1,523
5	Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94.	13.7	1,099
6	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. Nature Genetics, 2013, 45, 353-361.	9.4	960
7	Socioeconomic status and the 25â€^×â€^25 risk factors as determinants of premature mortality: a multicohort study and meta-analysis of 1Â∙7 million men and women. Lancet, The, 2017, 389, 1229-1237.	6.3	825
8	Multiple newly identified loci associated with prostate cancer susceptibility. Nature Genetics, 2008, 40, 316-321.	9.4	796
9	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. PLoS Medicine, 2010, 7, e1000279.	3.9	764
10	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. American Journal of Human Genetics, 2019, 104, 21-34.	2.6	711
11	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. Nature Genetics, 2018, 50, 928-936.	9.4	652
12	Iron-Overload–Related Disease in <i>HFE</i> Hereditary Hemochromatosis. New England Journal of Medicine, 2008, 358, 221-230.	13.9	649
13	Associations of Breast Cancer Risk Factors With Tumor Subtypes: A Pooled Analysis From the Breast Cancer Association Consortium Studies. Journal of the National Cancer Institute, 2011, 103, 250-263.	3.0	596
14	A common coding variant in CASP8 is associated with breast cancer risk. Nature Genetics, 2007, 39, 352-358.	9.4	591
15	Nonâ€melanoma skin cancer in Australia: the 2002 national survey and trends since 1985. Medical Journal of Australia, 2006, 184, 6-10.	0.8	559
16	Parent-of-origin-specific allelic associations among 106 genomic loci for age at menarche. Nature, 2014, 514, 92-97.	13.7	548
17	Heritability of Mammographic Density, a Risk Factor for Breast Cancer. New England Journal of Medicine, 2002, 347, 886-894.	13.9	537
18	The Anti Cancer Council of Victoria FFQ: relative validity of nutrient intakes compared with weighed food records in young to middleâ€aged women in a study of iron supplementation. Australian and New Zealand Journal of Public Health, 2000, 24, 576-583.	0.8	534

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19	Breast Cancer Risk Genes — Association Analysis in More than 113,000 Women. New England Journal of Medicine, 2021, 384, 428-439.	13.9	532
20	Detectable clonal mosaicism and its relationship to aging and cancer. Nature Genetics, 2012, 44, 651-658.	9.4	519
21	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
22	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. Nature Genetics, 2015, 47, 373-380.	9.4	513
23	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. Nature Genetics, 2013, 45, 371-384.	9.4	493
24	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. Nature Genetics, 2013, 45, 385-391.	9.4	492
25	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. Nature Genetics, 2009, 41, 585-590.	9.4	434
26	Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. Journal of the National Cancer Institute, 2015, 107, .	3.0	428
27	Genomic analyses identify hundreds of variants associated with age at menarche and support a role for puberty timing in cancer risk. Nature Genetics, 2017, 49, 834-841.	9.4	426
28	A novel recurrent mutation in MITF predisposes to familial and sporadic melanoma. Nature, 2011, 480, 99-103.	13.7	413
29	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. Nature Genetics, 2014, 46, 1103-1109.	9.4	408
30	Validation of Questionnaire and Bronchial Hyperresponsiveness against Respiratory Physician Assessment in the Diagnosis of Asthma. International Journal of Epidemiology, 1996, 25, 609-616.	0.9	397
31	Identification of seven new prostate cancer susceptibility loci through a genome-wide association study. Nature Genetics, 2009, 41, 1116-1121.	9.4	389
32	Childhood predictors of lung function trajectories and future COPD risk: a prospective cohort study from the first to the sixth decade of life. Lancet Respiratory Medicine,the, 2018, 6, 535-544.	5.2	381
33	Discovery of common and rare genetic risk variants for colorectal cancer. Nature Genetics, 2019, 51, 76-87.	9.4	377
34	Glycemic Index and Dietary Fiber and the Risk of Type 2 Diabetes. Diabetes Care, 2004, 27, 2701-2706.	4.3	374
35	Genome-wide association studies identify four ER negative–specific breast cancer risk loci. Nature Genetics, 2013, 45, 392-398.	9.4	374
36	Incidence of Malignant Disease in Biopsy-Proven Inflammatory Myopathy: A Population-Based Cohort Study. Annals of Internal Medicine, 2001, 134, 1087.	2.0	360

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37	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. Nature Genetics, 2015, 47, 1294-1303.	9.4	357
38	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. Nature Genetics, 2017, 49, 680-691.	9.4	356
39	Risk of cancer after use of fertility drugs with in-vitro fertilisation. Lancet, The, 1999, 354, 1586-1590.	6.3	349
40	Prevalence and Penetrance of Major Genes and Polygenes for Colorectal Cancer. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 404-412.	1.1	341
41	Risks of Lynch Syndrome Cancers for MSH6 Mutation Carriers. Journal of the National Cancer Institute, 2010, 102, 193-201.	3.0	328
42	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. Nature Genetics, 2013, 45, 362-370.	9.4	326
43	ï‰-3 Polyunsaturated Fatty Acid Biomarkers and Coronary Heart Disease. JAMA Internal Medicine, 2016, 176, 1155.	2.6	326
44	Incidence of non-melanocytic skin cancer treated in Australia. BMJ: British Medical Journal, 1988, 296, 13-17.	2.4	321
45	Heterogeneity of Breast Cancer Associations with Five Susceptibility Loci by Clinical and Pathological Characteristics. PLoS Genetics, 2008, 4, e1000054.	1.5	315
46	A Pooled Analysis of Waist Circumference and Mortality in 650,000 Adults. Mayo Clinic Proceedings, 2014, 89, 335-345.	1.4	307
47	Multiple Loci With Different Cancer Specificities Within the 8q24 Gene Desert. Journal of the National Cancer Institute, 2008, 100, 962-966.	3.0	306
48	Pathology Features in Bethesda Guidelines Predict Colorectal Cancer Microsatellite Instability: A Population-Based Study. Gastroenterology, 2007, 133, 48-56.	0.6	302
49	Association between Class III Obesity (BMI of 40–59 kg/m2) and Mortality: A Pooled Analysis of 20 Prospective Studies. PLoS Medicine, 2014, 11, e1001673.	3.9	299
50	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. Nature Genetics, 2014, 46, 994-1000.	9.4	294
51	Obesity and Outcomes in Premenopausal and Postmenopausal Breast Cancer. Cancer Epidemiology Biomarkers and Prevention, 2005, 14, 1686-1691.	1.1	290
52	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778.	9.4	289
53	Colorectal and Other Cancer Risks for Carriers and Noncarriers From Families With a DNA Mismatch Repair Gene Mutation: A Prospective Cohort Study. Journal of Clinical Oncology, 2012, 30, 958-964.	0.8	286
54	Breast Cancer Risk From Modifiable and Nonmodifiable Risk Factors Among White Women in the United States. JAMA Oncology, 2016, 2, 1295.	3.4	285

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55	A common variant at the TERT-CLPTM1L locus is associated with estrogen receptor–negative breast cancer. Nature Genetics, 2011, 43, 1210-1214.	9.4	279
56	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 126-135.	1.1	278
57	A genome-wide association study identifies a new ovarian cancer susceptibility locus on 9p22.2. Nature Genetics, 2009, 41, 996-1000.	9.4	276
58	Multiple loci on 8q24 associated with prostate cancer susceptibility. Nature Genetics, 2009, 41, 1058-1060.	9.4	273
59	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. Nature Genetics, 2011, 43, 785-791.	9.4	265
60	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. Nature Genetics, 2020, 52, 572-581.	9.4	265
61	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. Nature Genetics, 2021, 53, 65-75.	9.4	264
62	Insulin-like Growth Factors, Their Binding Proteins, and Prostate Cancer Risk: Analysis of Individual Patient Data from 12 Prospective Studies. Annals of Internal Medicine, 2008, 149, 461.	2.0	263
63	Mammographic Density Phenotypes and Risk of Breast Cancer: A Meta-analysis. Journal of the National Cancer Institute, 2014, 106, .	3.0	261
64	Genome-wide association analysis identifies three new breast cancer susceptibility loci. Nature Genetics, 2012, 44, 312-318.	9.4	256
65	Age at natural menopause and risk of incident cardiovascular disease: a pooled analysis of individual patient data. Lancet Public Health, The, 2019, 4, e553-e564.	4.7	252
66	Plasma phospholipid and dietary fatty acids as predictors of type 2 diabetes: interpreting the role of linoleic acid. American Journal of Clinical Nutrition, 2007, 86, 189-197.	2.2	251
67	Inositol polyphosphate 4-phosphatase II regulates PI3K/Akt signaling and is lost in human basal-like breast cancers. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 22231-22236.	3.3	249
68	The histologic phenotypes of breast carcinoma occurring before age 40 years in women with and without BRCA1 or BRCA2 germline mutations. Cancer, 1998, 83, 2335-2345.	2.0	243
69	Trends in the incidence of non-melanocytic skin cancer (NMSC) treated in Australia 1985–1995: Are primary prevention programs starting to have an effect?. , 1998, 78, 144-148.		241
70	Genome-wide association study identifies three new melanoma susceptibility loci. Nature Genetics, 2011, 43, 1108-1113.	9.4	230
71	Metachronous colorectal cancer risk for mismatch repair gene mutation carriers: the advantage of more extensive colon surgery. Gut, 2011, 60, 950-957.	6.1	227
72	Determining Risk of Colorectal Cancer and Starting Age of Screening Based on Lifestyle, Environmental, and Genetic Factors. Gastroenterology, 2018, 154, 2152-2164.e19.	0.6	226

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73	The INTERPHONE study: design, epidemiological methods, and description of the study population. European Journal of Epidemiology, 2007, 22, 647-664.	2.5	225
74	Genome-wide association study of glioma and meta-analysis. Human Genetics, 2012, 131, 1877-1888.	1.8	222
75	A pooled analysis of 14 cohort studies of anthropometric factors and pancreatic cancer risk. International Journal of Cancer, 2011, 129, 1708-1717.	2.3	221
76	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. Nature Genetics, 2015, 47, 164-171.	9.4	221
77	Omega-6 fatty acid biomarkers and incident type 2 diabetes: pooled analysis of individual-level data for 39†740 adults from 20 prospective cohort studies. Lancet Diabetes and Endocrinology,the, 2017, 5, 965-974.	5.5	213
78	Association of Body Mass Index and Age With Subsequent Breast Cancer Risk in Premenopausal Women. JAMA Oncology, 2018, 4, e181771.	3.4	210
79	Common sequence variants on 20q11.22 confer melanoma susceptibility. Nature Genetics, 2008, 40, 838-840.	9.4	209
80	Role of medical history in brain tumour development. Results from the international adult brain tumour study. , 1999, 82, 155-160.		205
81	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.	2.6	201
82	Cancer Risks for <i>MLH1</i> and <i>MSH2</i> Mutation Carriers. Human Mutation, 2013, 34, 490-497.	1.1	201
83	Biomarkers of Dietary Omega-6 Fatty Acids and Incident Cardiovascular Disease and Mortality. Circulation, 2019, 139, 2422-2436.	1.6	199
84	Hypomethylation of smoking-related genes is associated with future lung cancer in four prospective cohorts. Nature Communications, 2015, 6, 10192.	5.8	197
85	Use of Molecular Tumor Characteristics to Prioritize Mismatch Repair Gene Testing in Early-Onset Colorectal Cancer. Journal of Clinical Oncology, 2005, 23, 6524-6532.	0.8	194
86	Sunbed use during adolescence and early adulthood is associated with increased risk of earlyâ€onset melanoma. International Journal of Cancer, 2011, 128, 2425-2435.	2.3	194
87	Risks of Primary Extracolonic Cancers Following Colorectal Cancer in Lynch Syndrome. Journal of the National Cancer Institute, 2012, 104, 1363-1372.	3.0	193
88	Physical activity and risks of breast and colorectal cancer: a Mendelian randomisation analysis. Nature Communications, 2020, 11, 597.	5.8	193
89	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. Nature Communications, 2018, 9, 556.	5.8	188
90	Associations Between Dietary Nutrient Intake and Muscle Mass and Strength in Communityâ€Dwelling Older Adults: The Tasmanian Older Adult Cohort Study. Journal of the American Geriatrics Society, 2010, 58, 2129-2134.	1.3	184

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91	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. Nature Genetics, 2018, 50, 968-978.	9.4	184
92	Social adversity and epigenetic aging: a multi-cohort study on socioeconomic differences in peripheral blood DNA methylation. Scientific Reports, 2017, 7, 16266.	1.6	181
93	Risk of Colorectal Cancer for Carriers of Mutations in MUTYH, WithÂand Without a Family History of Cancer. Gastroenterology, 2014, 146, 1208-1211.e5.	0.6	180
94	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. Nature Genetics, 2013, 45, 868-876.	9.4	179
95	Identification of nine new susceptibility loci for endometrial cancer. Nature Communications, 2018, 9, 3166.	5.8	178
96	Incidence of melanoma and other malignancies among rheumatoid arthritis patients treated with methotrexate. Arthritis and Rheumatism, 2008, 59, 794-799.	6.7	172
97	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, 5373-5384.	1.4	168
98	HOXB13 is a susceptibility gene for prostate cancer: results from the International Consortium for Prostate Cancer Genetics (ICPCG). Human Genetics, 2013, 132, 5-14.	1.8	166
99	Fruit and Vegetable Intake and Risk of Breast Cancer by Hormone Receptor Status. Journal of the National Cancer Institute, 2013, 105, 219-236.	3.0	164
100	<i>CHEK2</i> *1100delC Heterozygosity in Women With Breast Cancer Associated With Early Death, Breast Cancer–Specific Death, and Increased Risk of a Second Breast Cancer. Journal of Clinical Oncology, 2012, 30, 4308-4316.	0.8	162
101	Genome-wide association study identifies new prostate cancer susceptibility loci. Human Molecular Genetics, 2011, 20, 3867-3875.	1.4	160
102	Circulating Steroid Hormones and the Risk of Prostate Cancer. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 86-91.	1.1	159
103	Alcohol Intake and Pancreatic Cancer Risk: A Pooled Analysis of Fourteen Cohort Studies. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 765-776.	1.1	158
104	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. Cancer Discovery, 2016, 6, 1052-1067.	7.7	157
105	BRCA2 Mutation-associated Breast Cancers Exhibit a Distinguishing Phenotype Based on Morphology and Molecular Profiles From Tissue Microarrays. American Journal of Surgical Pathology, 2007, 31, 121-128.	2.1	156
106	Adaptive evolution of the tumour suppressor BRCA1 in humans and chimpanzees. Nature Genetics, 2000, 25, 410-413.	9.4	153
107	DNA methylationâ€based biological aging and cancer risk and survival: Pooled analysis of seven prospective studies. International Journal of Cancer, 2018, 142, 1611-1619.	2.3	153
108	Low penetrance breast cancer susceptibility loci are associated with specific breast tumor subtypes: findings from the Breast Cancer Association Consortium. Human Molecular Genetics, 2011, 20, 3289-3303.	1.4	152

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109	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. Journal of the National Cancer Institute, 2015, 107, djv279.	3.0	152
110	Age- and Tumor Subtype–Specific Breast Cancer Risk Estimates for <i>CHEK2</i> *1100delC Carriers. Journal of Clinical Oncology, 2016, 34, 2750-2760.	0.8	152
111	Cancer Risks For Mismatch Repair Gene Mutation Carriers: A Population-Based Early Onset Case-Family Study. Clinical Gastroenterology and Hepatology, 2006, 4, 489-498.	2.4	151
112	Familial Risks, Early-Onset Breast Cancer, and BRCA1 and BRCA2 Germline Mutations. Journal of the National Cancer Institute, 2003, 95, 448-457.	3.0	150
113	Hormone replacement therapy and accuracy of mammographic screening. Lancet, The, 2000, 355, 270-274.	6.3	149
114	Multiple Novel Prostate Cancer Predisposition Loci Confirmed by an International Study: The PRACTICAL Consortium. Cancer Epidemiology Biomarkers and Prevention, 2008, 17, 2052-2061.	1.1	148
115	Anthropometric Factors and Thyroid Cancer Risk by Histological Subtype: Pooled Analysis of 22 Prospective Studies. Thyroid, 2016, 26, 306-318.	2.4	148
116	Constitutional Methylation of the <i>BRCA1</i> Promoter Is Specifically Associated with <i>BRCA1</i> Mutation-Associated Pathology in Early-Onset Breast Cancer. Cancer Prevention Research, 2011, 4, 23-33.	0.7	147
117	Genome-wide association study identifies multiple susceptibility loci for diffuse large B cell lymphoma. Nature Genetics, 2014, 46, 1233-1238.	9.4	147
118	Genome-wide association study identifies 25 known breast cancer susceptibility loci as risk factors for triple-negative breast cancer. Carcinogenesis, 2014, 35, 1012-1019.	1.3	145
119	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. Nature Communications, 2013, 4, 1628.	5.8	144
120	Fatty acid biomarkers of dairy fat consumption and incidence of type 2 diabetes: A pooled analysis of prospective cohort studies. PLoS Medicine, 2018, 15, e1002670.	3.9	143
121	Familial Patterns of Covariation for Cardiovascular Risk Factors in Adults: The Victorian Family Heart Study. American Journal of Epidemiology, 2000, 152, 704-715.	1.6	140
122	Genome-wide association study identifies a new melanoma susceptibility locus at 1q21.3. Nature Genetics, 2011, 43, 1114-1118.	9.4	140
123	A Combined Genomewide Linkage Scan of 1,233 Families for Prostate Cancer–Susceptibility Genes Conducted by the International Consortium for Prostate Cancer Genetics. American Journal of Human Genetics, 2005, 77, 219-229.	2.6	138
124	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. Nature Communications, 2015, 6, 7138.	5.8	138
125	Socioeconomic position, lifestyle habits and biomarkers of epigenetic aging: a multi-cohort analysis. Aging, 2019, 11, 2045-2070.	1.4	137
126	Evidence of Gene–Environment Interactions between Common Breast Cancer Susceptibility Loci and Established Environmental Risk Factors. PLoS Genetics, 2013, 9, e1003284.	1.5	136

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127	Analysis of cancer risk and BRCA1 and BRCA2mutation prevalence in the kConFab familial breast cancer resource. Breast Cancer Research, 2006, 8, R12.	2.2	135
128	Breast Cancer Prognosis in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers: An International Prospective Breast Cancer Family Registry Population-Based Cohort Study. Journal of Clinical Oncology, 2012, 30, 19-26.	0.8	134
129	Oral Contraceptive Use and Risk of Early-Onset Breast Cancer in Carriers and Noncarriers of BRCA1 and BRCA2 Mutations. Cancer Epidemiology Biomarkers and Prevention, 2005, 14, 350-356.	1.1	133
130	Plasma phospholipid fatty acid composition as a biomarker of habitual dietary fat intake in an ethnically diverse cohort. Nutrition, Metabolism and Cardiovascular Diseases, 2007, 17, 415-426.	1.1	133
131	Blood n-3 fatty acid levels and total and cause-specific mortality from 17 prospective studies. Nature Communications, 2021, 12, 2329.	5.8	132
132	Relationship between body adiposity measures and risk of primary knee and hip replacement for osteoarthritis: a prospective cohort study. Arthritis Research and Therapy, 2009, 11, R31.	1.6	131
133	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. Journal of the National Cancer Institute, 2019, 111, 146-157.	3.0	129
134	Expression of MUC2, MUC5AC, MUC5B, and MUC6 mucins in colorectal cancers and their association with the CpG island methylator phenotype. Modern Pathology, 2013, 26, 1642-1656.	2.9	127
135	The CYP3A4*1B polymorphism has no functional significance and is not associated with risk of breast or ovarian cancer. Pharmacogenetics and Genomics, 2002, 12, 355-366.	5.7	126
136	Colorectal carcinomas with KRAS mutation are associated with distinctive morphological and molecular features. Modern Pathology, 2013, 26, 825-834.	2.9	126
137	Sun exposure predicts risk of ocular melanoma in Australia. International Journal of Cancer, 2002, 101, 175-182.	2.3	125
138	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. Nature Genetics, 2016, 48, 374-386.	9.4	125
139	Genome-wide Modeling of Polygenic Risk Score in Colorectal Cancer Risk. American Journal of Human Genetics, 2020, 107, 432-444.	2.6	124
140	HFE C282Y homozygotes are at increased risk of breast and colorectal cancer. Hepatology, 2010, 51, 1311-1318.	3.6	123
141	The association between waist circumference and risk of mortality considering body mass index in 65- to 74-year-olds: a meta-analysis of 29 cohorts involving more than 58 000 elderly persons. International Journal of Epidemiology, 2012, 41, 805-817.	0.9	123
142	Early menarche, nulliparity and the risk for premature and early natural menopause. Human Reproduction, 2017, 32, 679-686.	0.4	122
143	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. Nature Genetics, 2020, 52, 56-73.	9.4	120
144	Recall bias in the assessment of exposure to mobile phones. Journal of Exposure Science and Environmental Epidemiology, 2009, 19, 369-381.	1.8	119

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145	A meta-analysis of genome-wide association studies to identify prostate cancer susceptibility loci associated with aggressive and non-aggressive disease. Human Molecular Genetics, 2013, 22, 408-415.	1.4	118
146	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. PLoS Medicine, 2016, 13, e1002105.	3.9	118
147	Foods, nutrients and prostate cancer. Cancer Causes and Control, 2004, 15, 11-20.	0.8	117
148	PIK3CA Activating Mutation in Colorectal Carcinoma: Associations with Molecular Features and Survival. PLoS ONE, 2013, 8, e65479.	1.1	117
149	A Meta-analysis of Individual Participant Data Reveals an Association between Circulating Levels of IGF-I and Prostate Cancer Risk. Cancer Research, 2016, 76, 2288-2300.	0.4	117
150	Traffic-related air pollution exposure is associated with allergic sensitization, asthma, and poor lung function in middle age. Journal of Allergy and Clinical Immunology, 2017, 139, 122-129.e1.	1.5	117
151	Fat Consumption and Its Association With Age-Related Macular Degeneration. JAMA Ophthalmology, 2009, 127, 674.	2.6	116
152	10-year performance of four models of breast cancer risk: a validation study. Lancet Oncology, The, 2019, 20, 504-517.	5.1	116
153	BRCA1 and BRCA2 Mutation Carriers, Oral Contraceptive Use, and Breast Cancer Before Age 50. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 1863-1870.	1.1	115
154	DNA methylation changes measured in preâ€diagnostic peripheral blood samples are associated with smoking and lung cancer risk. International Journal of Cancer, 2017, 140, 50-61.	2.3	115
155	Circulating steroid hormone concentrations in postmenopausal women in relation to body size and composition. Breast Cancer Research and Treatment, 2009, 115, 171-179.	1.1	113
156	Seasonality of cardiovascular risk factors: an analysis including over 230â€000 participants in 15 countries. Heart, 2014, 100, 1517-1523.	1.2	113
157	A Comparison of Adiposity Measures as Predictors of Allâ€cause Mortality: The Melbourne Collaborative Cohort Study. Obesity, 2007, 15, 994-1003.	1.5	112
158	A genome wide linkage search for breast cancer susceptibility genes. Genes Chromosomes and Cancer, 2006, 45, 646-655.	1.5	111
159	Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. International Journal of Epidemiology, 2016, 45, 1619-1630.	0.9	111
160	Childhood Lung Function Predicts Adult Chronic Obstructive Pulmonary Disease and Asthma–Chronic Obstructive Pulmonary Disease Overlap Syndrome. American Journal of Respiratory and Critical Care Medicine, 2017, 196, 39-46.	2.5	111
161	A case-control study of melanomas of the soles and palms (Australia and Scotland). Cancer Causes and Control, 1999, 10, 21-25.	0.8	110
162	Incidence of total knee and hip replacement for osteoarthritis in relation to the metabolic syndrome and its components: A prospective cohort study. Seminars in Arthritis and Rheumatism, 2014, 43, 429-436.	1.6	110

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163	Cumulative Burden of Colorectal Cancer–Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. Gastroenterology, 2020, 158, 1274-1286.e12.	0.6	110
164	The epidemiology of prostate cancer. Urologic Clinics of North America, 2003, 30, 209-217.	0.8	109
165	Common Breast Cancer Susceptibility Loci Are Associated with Triple-Negative Breast Cancer. Cancer Research, 2011, 71, 6240-6249.	0.4	109
166	Genome-wide association study identifies multiple loci associated with both mammographic density and breast cancer risk. Nature Communications, 2014, 5, 5303.	5.8	109
167	Effect of physical activity on articular knee joint structures in communityâ€based adults. Arthritis and Rheumatism, 2007, 57, 1261-1268.	6.7	108
168	The Interplay between the Effects of Lifetime Asthma, Smoking, and Atopy on Fixed Airflow Obstruction in Middle Age. American Journal of Respiratory and Critical Care Medicine, 2013, 187, 42-48.	2.5	108
169	Mammographic density and ageing: A collaborative pooled analysis of cross-sectional data from 22 countries worldwide. PLoS Medicine, 2017, 14, e1002335.	3.9	108
170	Dietary Patterns and Diabetes Incidence in the Melbourne Collaborative Cohort Study. American Journal of Epidemiology, 2007, 165, 603-610.	1.6	107
171	Carotenoids, retinol, tocopherols, and prostate cancer risk: pooled analysis of 15 studies. American Journal of Clinical Nutrition, 2015, 102, 1142-1157.	2.2	107
172	Combined genetic and splicing analysis of BRCA1 c.[594-2A>C; 641A>G] highlights the relevance of naturally occurring in-frame transcripts for developing disease gene variant classification algorithms. Human Molecular Genetics, 2016, 25, 2256-2268.	1.4	106
173	Association of DNA Methylation-Based Biological Age With Health Risk Factors and Overall and Cause-Specific Mortality. American Journal of Epidemiology, 2018, 187, 529-538.	1.6	106
174	Body size and composition and the risk of gastric and oesophageal adenocarcinoma. International Journal of Cancer, 2006, 118, 2628-2631.	2.3	105
175	Measurement of Serum Levels of Macrophage Inhibitory Cytokine 1 Combined with Prostate-Specific Antigen Improves Prostate Cancer Diagnosis. Clinical Cancer Research, 2006, 12, 89-96.	3.2	105
176	Identification of a BRCA2-Specific Modifier Locus at 6p24 Related to Breast Cancer Risk. PLoS Genetics, 2013, 9, e1003173.	1.5	105
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