

Christopher A Haiman

List of Publications by Year in descending order

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Version: 2024-02-01

590
papers

57,917
citations

2203

99
h-index

1820

210
g-index

610
all docs

610
docs citations

610
times ranked

57675
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015, 518, 197-206.	13.7	3,823
2	Age-Related Clonal Hematopoiesis Associated with Adverse Outcomes. <i>New England Journal of Medicine</i> , 2014, 371, 2488-2498.	13.9	3,474
3	Genome-wide association study identifies novel breast cancer susceptibility loci. <i>Nature</i> , 2007, 447, 1087-1093.	13.7	2,165
4	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014, 46, 1173-1186.	9.4	1,818
5	Risk of COVID-19 among front-line health-care workers and the general community: a prospective cohort study. <i>Lancet Public Health</i> , The, 2020, 5, e475-e483.	4.7	1,595
6	New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015, 518, 187-196.	13.7	1,328
7	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	13.7	1,099
8	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	9.4	960
9	Common variants on chromosomes 2q35 and 16q12 confer susceptibility to estrogen receptor-positive breast cancer. <i>Nature Genetics</i> , 2007, 39, 865-869.	9.4	774
10	Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. <i>American Journal of Human Genetics</i> , 2019, 104, 21-34.	2.6	711
11	Genetic analyses of diverse populations improves discovery for complex traits. <i>Nature</i> , 2019, 570, 514-518.	13.7	679
12	Ethnic and Racial Differences in the Smoking-Related Risk of Lung Cancer. <i>New England Journal of Medicine</i> , 2006, 354, 333-342.	13.9	668
13	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. <i>Nature Genetics</i> , 2018, 50, 928-936.	9.4	652
14	Multiple regions within 8q24 independently affect risk for prostate cancer. <i>Nature Genetics</i> , 2007, 39, 638-644.	9.4	621
15	The 8q24 cancer risk variant rs6983267 shows long-range interaction with MYC in colorectal cancer. <i>Nature Genetics</i> , 2009, 41, 882-884.	9.4	616
16	Associations of Breast Cancer Risk Factors With Tumor Subtypes: A Pooled Analysis From the Breast Cancer Association Consortium Studies. <i>Journal of the National Cancer Institute</i> , 2011, 103, 250-263.	3.0	596
17	Admixture mapping identifies 8q24 as a prostate cancer risk locus in African-American men. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 14068-14073.	3.3	575
18	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , 2012, 44, 651-658.	9.4	519

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19	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015, 47, 373-380.	9.4	513
20	Detectable clonal mosaicism from birth to old age and its relationship to cancer. <i>Nature Genetics</i> , 2012, 44, 642-650.	9.4	511
21	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384.	9.4	493
22	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. <i>Nature Genetics</i> , 2013, 45, 385-391.	9.4	492
23	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. <i>Nature Genetics</i> , 2017, 49, 1126-1132.	9.4	472
24	Sequence variants in SLC16A11 are a common risk factor for type 2 diabetes in Mexico. <i>Nature</i> , 2014, 506, 97-101.	13.7	439
25	A Population-Based Study of Genes Previously Implicated in Breast Cancer. <i>New England Journal of Medicine</i> , 2021, 384, 440-451.	13.9	414
26	Common variants on chromosome 5p12 confer susceptibility to estrogen receptor- α positive breast cancer. <i>Nature Genetics</i> , 2008, 40, 703-706.	9.4	412
27	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. <i>Nature Genetics</i> , 2014, 46, 1103-1109.	9.4	408
28	Identification of seven new prostate cancer susceptibility loci through a genome-wide association study. <i>Nature Genetics</i> , 2009, 41, 1116-1121.	9.4	389
29	Choosing Haplotype-Tagging SNPS Based on Unphased Genotype Data Using a Preliminary Sample of Unrelated Subjects with an Example from the Multiethnic Cohort Study. <i>Human Heredity</i> , 2003, 55, 27-36.	0.4	386
30	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	9.4	374
31	Meta-analysis identifies common variants associated with body mass index in east Asians. <i>Nature Genetics</i> , 2012, 44, 307-311.	9.4	372
32	Large-scale genomic analyses link reproductive aging to hypothalamic signaling, breast cancer susceptibility and BRCA1-mediated DNA repair. <i>Nature Genetics</i> , 2015, 47, 1294-1303.	9.4	357
33	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691.	9.4	356
34	A common genetic risk factor for colorectal and prostate cancer. <i>Nature Genetics</i> , 2007, 39, 954-956.	9.4	336
35	The landscape of recombination in African Americans. <i>Nature</i> , 2011, 476, 170-175.	13.7	319
36	Heterogeneity of Breast Cancer Associations with Five Susceptibility Loci by Clinical and Pathological Characteristics. <i>PLoS Genetics</i> , 2008, 4, e1000054.	1.5	315

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37	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. <i>Nature Genetics</i> , 2014, 46, 994-1000.	9.4	294
38	Genome-Wide Association Study of Coronary Heart Disease and Its Risk Factors in 8,090 African Americans: The NHLBI CARE Project. <i>PLoS Genetics</i> , 2011, 7, e1001300.	1.5	290
39	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778.	9.4	289
40	Breast Cancer Risk From Modifiable and Nonmodifiable Risk Factors Among White Women in the United States. <i>JAMA Oncology</i> , 2016, 2, 1295.	3.4	285
41	A common variant at the TERT-CLPTM1L locus is associated with estrogen receptor-negative breast cancer. <i>Nature Genetics</i> , 2011, 43, 1210-1214.	9.4	279
42	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 126-135.	1.1	278
43	A Genomewide Admixture Map for Latino Populations. <i>American Journal of Human Genetics</i> , 2007, 80, 1024-1036.	2.6	265
44	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. <i>Nature Genetics</i> , 2011, 43, 785-791.	9.4	265
45	Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. <i>Nature Genetics</i> , 2020, 52, 572-581.	9.4	265
46	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. <i>Nature Genetics</i> , 2021, 53, 65-75.	9.4	264
47	Multi-ancestry genetic study of type 2 diabetes highlights the power of diverse populations for discovery and translation. <i>Nature Genetics</i> , 2022, 54, 560-572.	9.4	250
48	Exome sequencing of 20,791 cases of type 2 diabetes and 24,440 controls. <i>Nature</i> , 2019, 570, 71-76.	13.7	248
49	Generalization and Dilution of Association Results from European GWAS in Populations of Non-European Ancestry: The PAGE Study. <i>PLoS Biology</i> , 2013, 11, e1001661.	2.6	235
50	A meta-analysis identifies new loci associated with body mass index in individuals of African ancestry. <i>Nature Genetics</i> , 2013, 45, 690-696.	9.4	232
51	Association of a Low-Frequency Variant in <i>HNF1A</i> With Type 2 Diabetes in a Latino Population. <i>JAMA - Journal of the American Medical Association</i> , 2014, 311, 2305.	3.8	230
52	Transferability of tag SNPs in genetic association studies in multiple populations. <i>Nature Genetics</i> , 2006, 38, 1298-1303.	9.4	224
53	Identification of a new prostate cancer susceptibility locus on chromosome 8q24. <i>Nature Genetics</i> , 2009, 41, 1055-1057.	9.4	218
54	Functional Enhancers at the Gene-Poor 8q24 Cancer-Linked Locus. <i>PLoS Genetics</i> , 2009, 5, e1000597.	1.5	214

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55	Functional Variants at the 11q13 Risk Locus for Breast Cancer Regulate Cyclin D1 Expression through Long-Range Enhancers. <i>American Journal of Human Genetics</i> , 2013, 92, 489-503.	2.6	201
56	Circulating Vitamin D and Colorectal Cancer Risk: An International Pooling Project of 17 Cohorts. <i>Journal of the National Cancer Institute</i> , 2019, 111, 158-169.	3.0	199
57	Genome-wide association study of prostate cancer in men of African ancestry identifies a susceptibility locus at 17q21. <i>Nature Genetics</i> , 2011, 43, 570-573.	9.4	198
58	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , 2018, 9, 556.	5.8	188
59	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , 2018, 50, 968-978.	9.4	184
60	Genetic insights into biological mechanisms governing human ovarian ageing. <i>Nature</i> , 2021, 596, 393-397.	13.7	183
61	Association of Coffee Consumption With Total and Cause-Specific Mortality Among Nonwhite Populations. <i>Annals of Internal Medicine</i> , 2017, 167, 228.	2.0	182
62	Global Patterns of Prostate Cancer Incidence, Aggressiveness, and Mortality in Men of African Descent. <i>Prostate Cancer</i> , 2013, 2013, 1-12.	0.4	180
63	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018, 9, 3166.	5.8	178
64	<i>PALB2</i> , <i>CHEK2</i> and <i>ATM</i> rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016, 53, 800-811.	1.5	174
65	Phenome-Wide Association Study (PheWAS) for Detection of Pleiotropy within the Population Architecture using Genomics and Epidemiology (PAGE) Network. <i>PLoS Genetics</i> , 2013, 9, e1003087.	1.5	171
66	Genome-wide meta-analysis of 241,258 adults accounting for smoking behaviour identifies novel loci for obesity traits. <i>Nature Communications</i> , 2017, 8, 14977.	5.8	169
67	Consistent Association of Type 2 Diabetes Risk Variants Found in Europeans in Diverse Racial and Ethnic Groups. <i>PLoS Genetics</i> , 2010, 6, e1001078.	1.5	168
68	A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. <i>Human Molecular Genetics</i> , 2012, 21, 5373-5384.	1.4	168
69	Comprehensive Functional Annotation of 77 Prostate Cancer Risk Loci. <i>PLoS Genetics</i> , 2014, 10, e1004102.	1.5	167
70	The Next PAGE in Understanding Complex Traits: Design for the Analysis of Population Architecture Using Genetics and Epidemiology (PAGE) Study. <i>American Journal of Epidemiology</i> , 2011, 174, 849-859.	1.6	161
71	Genome-wide association study identifies new prostate cancer susceptibility loci. <i>Human Molecular Genetics</i> , 2011, 20, 3867-3875.	1.4	160
72	Genome-wide physical activity interactions in adiposity • A meta-analysis of 200,452 adults. <i>PLoS Genetics</i> , 2017, 13, e1006528.	1.5	158

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73	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. <i>Cancer Discovery</i> , 2016, 6, 1052-1067.	7.7	157
74	Low penetrance breast cancer susceptibility loci are associated with specific breast tumor subtypes: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2011, 20, 3289-3303.	1.4	152
75	Interactions Between Genetic Variants and Breast Cancer Risk Factors in the Breast and Prostate Cancer Cohort Consortium. <i>Journal of the National Cancer Institute</i> , 2011, 103, 1252-1263.	3.0	147
76	Genetic Determinants of Lipid Traits in Diverse Populations from the Population Architecture using Genomics and Epidemiology (PAGE) Study. <i>PLoS Genetics</i> , 2011, 7, e1002138.	1.5	146
77	A comprehensive haplotype analysis of CYP19 and breast cancer risk: the Multiethnic Cohort. <i>Human Molecular Genetics</i> , 2003, 12, 2679-2692.	1.4	144
78	Genetic Variation at the CYP19A1 Locus Predicts Circulating Estrogen Levels but not Breast Cancer Risk in Postmenopausal Women. <i>Cancer Research</i> , 2007, 67, 1893-1897.	0.4	140
79	The gene, environment association studies consortium (GENEVA): maximizing the knowledge obtained from GWAS by collaboration across studies of multiple conditions. <i>Genetic Epidemiology</i> , 2010, 34, 364-372.	0.6	139
80	The contribution of rare variation to prostate cancer heritability. <i>Nature Genetics</i> , 2016, 48, 30-35.	9.4	139
81	Genome-Wide Association Study in East Asians Identifies Novel Susceptibility Loci for Breast Cancer. <i>PLoS Genetics</i> , 2012, 8, e1002532.	1.5	137
82	A Common 8q24 Variant in Prostate and Breast Cancer from a Large Nested Case-Control Study. <i>Cancer Research</i> , 2007, 67, 2951-2956.	0.4	136
83	Association of Diabetes With Prostate Cancer Risk in the Multiethnic Cohort. <i>American Journal of Epidemiology</i> , 2009, 169, 937-945.	1.6	136
84	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 146-157.	3.0	129
85	Genetic determinants of telomere length and risk of common cancers: a Mendelian randomization study. <i>Human Molecular Genetics</i> , 2015, 24, 5356-5366.	1.4	128
86	A tetranucleotide repeat polymorphism in CYP19 and breast cancer risk. <i>International Journal of Cancer</i> , 2000, 87, 204-210.	2.3	127
87	Breast cancer risk variants at 6q25 display different phenotype associations and regulate ESR1, RMND1 and CCDC170. <i>Nature Genetics</i> , 2016, 48, 374-386.	9.4	125
88	Mendelian randomization study of adiposity-related traits and risk of breast, ovarian, prostate, lung and colorectal cancer. <i>International Journal of Epidemiology</i> , 2016, 45, 896-908.	0.9	124
89	Cyclin D1b variant influences prostate cancer growth through aberrant androgen receptor regulation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 2190-2195.	3.3	123
90	Genome-wide association study of breast cancer in Latinas identifies novel protective variants on 6q25. <i>Nature Communications</i> , 2014, 5, 5260.	5.8	123

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91	Large-scale transcriptome-wide association study identifies new prostate cancer risk regions. <i>Nature Communications</i> , 2018, 9, 4079.	5.8	121
92	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	9.4	120
93	A meta-analysis of genome-wide association studies to identify prostate cancer susceptibility loci associated with aggressive and non-aggressive disease. <i>Human Molecular Genetics</i> , 2013, 22, 408-415.	1.4	118
94	Obesity, body fat distribution, and risk of breast cancer subtypes in African American women participating in the AMBER Consortium. <i>Breast Cancer Research and Treatment</i> , 2015, 150, 655-666.	1.1	118
95	Genetically Predicted Body Mass Index and Breast Cancer Risk: Mendelian Randomization Analyses of Data from 145,000 Women of European Descent. <i>PLoS Medicine</i> , 2016, 13, e1002105.	3.9	118
96	Characterizing Genetic Risk at Known Prostate Cancer Susceptibility Loci in African Americans. <i>PLoS Genetics</i> , 2011, 7, e1001387.	1.5	117
97	Trans-Ethnic Fine-Mapping of Lipid Loci Identifies Population-Specific Signals and Allelic Heterogeneity That Increases the Trait Variance Explained. <i>PLoS Genetics</i> , 2013, 9, e1003379.	1.5	112
98	A Large Multiethnic Genome-Wide Association Study of Prostate Cancer Identifies Novel Risk Variants and Substantial Ethnic Differences. <i>Cancer Discovery</i> , 2015, 5, 878-891.	7.7	111
99	Prostate Cancer Susceptibility in Men of African Ancestry at 8q24. <i>Journal of the National Cancer Institute</i> , 2016, 108, djv431.	3.0	111
100	FGFR2 variants and breast cancer risk: fine-scale mapping using African American studies and analysis of chromatin conformation. <i>Human Molecular Genetics</i> , 2009, 18, 1692-1703.	1.4	110
101	Enhanced Statistical Tests for GWAS in Admixed Populations: Assessment using African Americans from CARE and a Breast Cancer Consortium. <i>PLoS Genetics</i> , 2011, 7, e1001371.	1.5	110
102	Cumulative Burden of Colorectal Cancer-Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. <i>Gastroenterology</i> , 2020, 158, 1274-1286.e12.	0.6	110
103	Genome-Wide Association of Body Fat Distribution in African Ancestry Populations Suggests New Loci. <i>PLoS Genetics</i> , 2013, 9, e1003681.	1.5	109
104	Identification of a Functional Genetic Variant at 16q12.1 for Breast Cancer Risk: Results from the Asia Breast Cancer Consortium. <i>PLoS Genetics</i> , 2010, 6, e1001002.	1.5	107
105	IGF-1, IGFBP-1, and IGFBP-3 Polymorphisms Predict Circulating IGF Levels but Not Breast Cancer Risk: Findings from the Breast and Prostate Cancer Cohort Consortium (BPC3). <i>PLoS ONE</i> , 2008, 3, e2578.	1.1	106
106	Identification of a BRCA2-Specific Modifier Locus at 6p24 Related to Breast Cancer Risk. <i>PLoS Genetics</i> , 2013, 9, e1003173.	1.5	105
107	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	5.8	105
108	Generalizability of Associations from Prostate Cancer Genome-Wide Association Studies in Multiple Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 1285-1289.	1.1	102

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109	Common Breast Cancer Susceptibility Variants in <i>LSP1</i> and <i>RAD51L1</i> Are Associated with Mammographic Density Measures that Predict Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 1156-1166.	1.1	101
110	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	2.6	101
111	Fine-mapping identifies multiple prostate cancer risk loci at 5p15, one of which associates with TERT expression. <i>Human Molecular Genetics</i> , 2013, 22, 2520-2528.	1.4	100
112	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. <i>Cancer Research</i> , 2016, 76, 5103-5114.	0.4	100
113	Common variants at 11q12, 10q26 and 3p11.2 are associated with prostate cancer susceptibility in Japanese. <i>Nature Genetics</i> , 2012, 44, 426-429.	9.4	98
114	Fine-Scale Mapping of the FGFR2 Breast Cancer Risk Locus: Putative Functional Variants Differentially Bind FOXA1 and E2F1. <i>American Journal of Human Genetics</i> , 2013, 93, 1046-1060.	2.6	98
115	High-Quality Diets Associate With Reduced Risk of Colorectal Cancer: Analyses of Diet Quality Indexes in the Multiethnic Cohort. <i>Gastroenterology</i> , 2017, 153, 386-394.e2.	0.6	98
116	Discovery and fine-mapping of adiposity loci using high density imputation of genome-wide association studies in individuals of African ancestry: African Ancestry Anthropometry Genetics Consortium. <i>PLoS Genetics</i> , 2017, 13, e1006719.	1.5	98
117	A Review of Cancer in U.S. Hispanic Populations. <i>Cancer Prevention Research</i> , 2012, 5, 150-163.	0.7	95
118	No evidence that protein truncating variants in <i>BRIP1</i> are associated with breast cancer risk: implications for gene panel testing. <i>Journal of Medical Genetics</i> , 2016, 53, 298-309.	1.5	94
119	Identification, Replication, and Fine-Mapping of Loci Associated with Adult Height in Individuals of African Ancestry. <i>PLoS Genetics</i> , 2011, 7, e1002298.	1.5	93
120	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016, 7, 11375.	5.8	93
121	Genome-wide association study identifies breast cancer risk variant at 10q21.2: results from the Asia Breast Cancer Consortium. <i>Human Molecular Genetics</i> , 2011, 20, 4991-4999.	1.4	92
122	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014, 23, 6616-6633.	1.4	90
123	CAUSEL: an epigenome- and genome-editing pipeline for establishing function of noncoding GWAS variants. <i>Nature Medicine</i> , 2015, 21, 1357-1363.	15.2	90
124	Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. <i>Nature Communications</i> , 2019, 10, 1741.	5.8	90
125	Validation of Genome-Wide Prostate Cancer Associations in Men of African Descent. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 23-32.	1.1	88
126	Two susceptibility loci identified for prostate cancer aggressiveness. <i>Nature Communications</i> , 2015, 6, 6889.	5.8	88

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127	Three new pancreatic cancer susceptibility signals identified on chromosomes 1q32.1, 5p15.33 and 8q24.21. <i>Oncotarget</i> , 2016, 7, 66328-66343.	0.8	88
128	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. <i>Nature Communications</i> , 2018, 9, 2256.	5.8	88
129	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	5.8	88
130	Common genetic determinants of breast-cancer risk in East Asian women: a collaborative study of 23 637 breast cancer cases and 25 579 controls. <i>Human Molecular Genetics</i> , 2013, 22, 2539-2550.	1.4	86
131	A sequence repeat in the insulin-like growth factor-1 gene and risk of breast cancer. <i>International Journal of Cancer</i> , 2002, 100, 332-336.	2.3	84
132	Association of ESR1 gene tagging SNPs with breast cancer risk. <i>Human Molecular Genetics</i> , 2009, 18, 1131-1139.	1.4	84
133	Cyclin D1 Splice Variants: Polymorphism, Risk, and Isoform-Specific Regulation in Prostate Cancer. <i>Clinical Cancer Research</i> , 2009, 15, 5338-5349.	3.2	84
134	Associations of autozygosity with a broad range of human phenotypes. <i>Nature Communications</i> , 2019, 10, 4957.	5.8	84
135	Meta-analysis of up to 622,409 individuals identifies 40 novel smoking behaviour associated genetic loci. <i>Molecular Psychiatry</i> , 2020, 25, 2392-2409.	4.1	83
136	Fine mapping of the association with obesity at the FTO locus in African-derived populations. <i>Human Molecular Genetics</i> , 2010, 19, 2907-2916.	1.4	82
137	Racial/Ethnic Differences in Postmenopausal Endogenous Hormones: The Multiethnic Cohort Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 1849-1855.	1.1	79
138	Polymorphisms in Steroid Hormone Pathway Genes and Mammographic Density. <i>Breast Cancer Research and Treatment</i> , 2003, 77, 27-36.	1.1	78
139	Informed Conditioning on Clinical Covariates Increases Power in Case-Control Association Studies. <i>PLoS Genetics</i> , 2012, 8, e1003032.	1.5	78
140	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus. <i>Nature Communications</i> , 2016, 7, 12675.	5.8	78
141	Admixture Mapping of 15,280 African Americans Identifies Obesity Susceptibility Loci on Chromosomes 5 and X. <i>PLoS Genetics</i> , 2009, 5, e1000490.	1.5	78
142	Fine Mapping and Identification of BMI Loci in African Americans. <i>American Journal of Human Genetics</i> , 2013, 93, 661-671.	2.6	77
143	Genetic Variants Associated With Quantitative Glucose Homeostasis Traits Translate to Type 2 Diabetes in Mexican Americans: The GUARDIAN (Genetics Underlying Diabetes in Hispanics) Consortium. <i>Diabetes</i> , 2015, 64, 1853-1866.	0.3	77
144	The effects of height and BMI on prostate cancer incidence and mortality: a Mendelian randomization study in 20,848 cases and 20,214 controls from the PRACTICAL consortium. <i>Cancer Causes and Control</i> , 2015, 26, 1603-1616.	0.8	77

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145	Fine-Scale Mapping of the 5q11.2 Breast Cancer Locus Reveals at Least Three Independent Risk Variants Regulating MAP3K1. <i>American Journal of Human Genetics</i> , 2015, 96, 5-20.	2.6	76
146	Genetic and functional analysis of <i>CHEK2</i> (<i>CHK2</i>) variants in multiethnic cohorts. <i>International Journal of Cancer</i> , 2007, 121, 2661-2667.	2.3	75
147	<i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017, 77, 2789-2799.	0.4	75
148	Assessment of polygenic architecture and risk prediction based on common variants across fourteen cancers. <i>Nature Communications</i> , 2020, 11, 3353.	5.8	75
149	Racial/Ethnic Differences in Lung Cancer Incidence in the Multiethnic Cohort Study: An Update. <i>Journal of the National Cancer Institute</i> , 2019, 111, 811-819.	3.0	74
150	Generalizability and Epidemiologic Characterization of Eleven Colorectal Cancer GWAS Hits in Multiple Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 70-81.	1.1	73
151	The Dietary Inflammatory Index Is Associated with Colorectal Cancer Risk in the Multiethnic Cohort. <i>Journal of Nutrition</i> , 2017, 147, jn242529.	1.3	73
152	Trans-ethnic genome-wide association study of colorectal cancer identifies a new susceptibility locus in VT11A. <i>Nature Communications</i> , 2014, 5, 4613.	5.8	72
153	A genome-wide association study of prostate cancer in West African men. <i>Human Genetics</i> , 2014, 133, 509-521.	1.8	72
154	Strategies for Enriching Variant Coverage in Candidate Disease Loci on a Multiethnic Genotyping Array. <i>PLoS ONE</i> , 2016, 11, e0167758.	1.1	72
155	Replication and Functional Genomic Analyses of the Breast Cancer Susceptibility Locus at 6q25.1 Generalize Its Importance in Women of Chinese, Japanese, and European Ancestry. <i>Cancer Research</i> , 2011, 71, 1344-1355.	0.4	71
156	Evaluation of the Metabochip Genotyping Array in African Americans and Implications for Fine Mapping of GWAS-Identified Loci: The PAGE Study. <i>PLoS ONE</i> , 2012, 7, e35651.	1.1	71
157	Expression QTL-based analyses reveal candidate causal genes and loci across five tumor types. <i>Human Molecular Genetics</i> , 2014, 23, 5294-5302.	1.4	71
158	Dietary Red and Processed Meat Intake and Markers of Adiposity and Inflammation: The Multiethnic Cohort Study. <i>Journal of the American College of Nutrition</i> , 2017, 36, 378-385.	1.1	71
159	African Ancestry and Its Correlation to Type 2 Diabetes in African Americans: A Genetic Admixture Analysis in Three U.S. Population Cohorts. <i>PLoS ONE</i> , 2012, 7, e32840.	1.1	70
160	A genome-wide association study of breast cancer in women of African ancestry. <i>Human Genetics</i> , 2013, 132, 39-48.	1.8	70
161	Leveraging population admixture to characterize the heritability of complex traits. <i>Nature Genetics</i> , 2014, 46, 1356-1362.	9.4	69
162	Genome-wide Scan of 29,141 African Americans Finds No Evidence of Directional Selection since Admixture. <i>American Journal of Human Genetics</i> , 2014, 95, 437-444.	2.6	69

#	ARTICLE	IF	CITATIONS
163	Exome Chip Meta-analysis Fine Maps Causal Variants and Elucidates the Genetic Architecture of Rare Coding Variants in Smoking and Alcohol Use. <i>Biological Psychiatry</i> , 2019, 85, 946-955.	0.7	69
164	Association between Adult Height and Risk of Colorectal, Lung, and Prostate Cancer: Results from Meta-analyses of Prospective Studies and Mendelian Randomization Analyses. <i>PLoS Medicine</i> , 2016, 13, e1002118.	3.9	69
165	PTGS2 and IL6 genetic variation and risk of breast and prostate cancer: results from the Breast and Prostate Cancer Cohort Consortium (BPC3). <i>Carcinogenesis</i> , 2010, 31, 455-461.	1.3	68
166	Fine scale mapping of the breast cancer 16q12 locus. <i>Human Molecular Genetics</i> , 2010, 19, 2507-2515.	1.4	68
167	Blood lipids and prostate cancer: a Mendelian randomization analysis. <i>Cancer Medicine</i> , 2016, 5, 1125-1136.	1.3	68
168	Association of Type 2 Diabetes Susceptibility Variants With Advanced Prostate Cancer Risk in the Breast and Prostate Cancer Cohort Consortium. <i>American Journal of Epidemiology</i> , 2012, 176, 1121-1129.	1.6	67
169	Genome-Wide Testing of Putative Functional Exonic Variants in Relationship with Breast and Prostate Cancer Risk in a Multiethnic Population. <i>PLoS Genetics</i> , 2013, 9, e1003419.	1.5	67
170	Multiple novel prostate cancer susceptibility signals identified by fine-mapping of known risk loci among Europeans. <i>Human Molecular Genetics</i> , 2015, 24, 5589-5602.	1.4	67
171	The androgen receptor CAG repeat polymorphism and risk of breast cancer in the Nurses' Health Study. <i>Cancer Research</i> , 2002, 62, 1045-9.	0.4	67
172	Genetic Variation in the HSD17B1 Gene and Risk of Prostate Cancer. <i>PLoS Genetics</i> , 2005, 1, e68.	1.5	66
173	Genetic determinants of mammographic density. <i>Breast Cancer Research</i> , 2002, 4, R5.	2.2	65
174	Common genetic variation in IGF1, IGFBP-1, and IGFBP-3 in relation to mammographic density: a cross-sectional study. <i>Breast Cancer Research</i> , 2007, 9, R18.	2.2	65
175	Haplotype Analysis of the HSD17B1 Gene and Risk of Breast Cancer: A Comprehensive Approach to Multicenter Analyses of Prospective Cohort Studies. <i>Cancer Research</i> , 2006, 66, 2468-2475.	0.4	64
176	Sex and ethnic/racial-specific risk factors for gallbladder disease. <i>BMC Gastroenterology</i> , 2017, 17, 153.	0.8	64
177	Common Genetic Variation in the Sex Steroid Hormone-Binding Globulin (SHBG) Gene and Circulating SHBG Levels among Postmenopausal Women: The Multiethnic Cohort. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2005, 90, 2198-2204.	1.8	63
178	A Systematic Mapping Approach of 16q12.2/FTO and BMI in More Than 20,000 African Americans Narrows in on the Underlying Functional Variation: Results from the Population Architecture using Genomics and Epidemiology (PAGE) Study. <i>PLoS Genetics</i> , 2013, 9, e1003171.	1.5	63
179	Cross Cancer Genomic Investigation of Inflammation Pathway for Five Common Cancers: Lung, Ovary, Prostate, Breast, and Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv246.	3.0	63
180	Generalizability of established prostate cancer risk variants in men of African ancestry. <i>International Journal of Cancer</i> , 2015, 136, 1210-1217.	2.3	62

#	ARTICLE	IF	CITATIONS
181	Two Estrogen-Related Variants in <i>CYP19A1</i> and Endometrial Cancer Risk: A Pooled Analysis in the Epidemiology of Endometrial Cancer Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 242-247.	1.1	61
182	Fine-mapping of breast cancer susceptibility loci characterizes genetic risk in African Americans. <i>Human Molecular Genetics</i> , 2011, 20, 4491-4503.	1.4	61
183	Association of Methylenetetrahydrofolate Reductase Gene Polymorphisms and Sex-Specific Survival in Patients With Metastatic Colon Cancer. <i>Journal of Clinical Oncology</i> , 2007, 25, 3726-3731.	0.8	60
184	Admixture mapping identifies a locus on 6q25 associated with breast cancer risk in US Latinas. <i>Human Molecular Genetics</i> , 2012, 21, 1907-1917.	1.4	60
185	Identification of susceptibility pathways for the role of chromosome 15q25.1 in modifying lung cancer risk. <i>Nature Communications</i> , 2018, 9, 3221.	5.8	60
186	Eighteen Insulin-like Growth Factor Pathway Genes, Circulating Levels of IGF-I and Its Binding Protein, and Risk of Prostate and Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2877-2887.	1.1	59
187	Association of the FTO Obesity Risk Variant rs8050136 With Percentage of Energy Intake From Fat in Multiple Racial/Ethnic Populations. <i>American Journal of Epidemiology</i> , 2013, 178, 780-790.	1.6	59
188	Evidence that the 5p12 Variant rs10941679 Confers Susceptibility to Estrogen-Receptor-Positive Breast Cancer through FGF10 and MRPS30 Regulation. <i>American Journal of Human Genetics</i> , 2016, 99, 903-911.	2.6	59
189	A Transcriptome-Wide Association Study Identifies Novel Candidate Susceptibility Genes for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , 2020, 112, 1003-1012.	3.0	59
190	Prediction of breast cancer risk by genetic risk factors, overall and by hormone receptor status. <i>Journal of Medical Genetics</i> , 2012, 49, 601-608.	1.5	58
191	Is high vitamin B12 status a cause of lung cancer?. <i>International Journal of Cancer</i> , 2019, 145, 1499-1503.	2.3	58
192	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	3.0	57
193	Genome-Wide Association Studies of Cancer in Diverse Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 405-417.	1.1	57
194	Characterizing Associations and SNP-Environment Interactions for GWAS-Identified Prostate Cancer Risk Markers—Results from BPC3. <i>PLoS ONE</i> , 2011, 6, e17142.	1.1	57
195	Genetic variation in angiotensin I-converting enzyme (ACE) and breast cancer risk: the multiethnic cohort. <i>Cancer Research</i> , 2003, 63, 6984-7.	0.4	57
196	A Large-Scale Analysis of Genetic Variants within Putative miRNA Binding Sites in Prostate Cancer. <i>Cancer Discovery</i> , 2015, 5, 368-379.	7.7	56
197	Identification of Novel Genetic Markers of Breast Cancer Survival. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	3.0	56
198	Risk Analysis of Prostate Cancer in PRACTICAL, a Multinational Consortium, Using 25 Known Prostate Cancer Susceptibility Loci. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1121-1129.	1.1	56

#	ARTICLE	IF	CITATIONS
199	Mercapturic Acids Derived from the Toxicants Acrolein and Crotonaldehyde in the Urine of Cigarette Smokers from Five Ethnic Groups with Differing Risks for Lung Cancer. <i>PLoS ONE</i> , 2015, 10, e0124841.	1.1	56
200	Novel Associations between Common Breast Cancer Susceptibility Variants and Risk-Predicting Mammographic Density Measures. <i>Cancer Research</i> , 2015, 75, 2457-2467.	0.4	55
201	<i>A priori</i>-defined diet quality indices, biomarkers and risk for type 2 diabetes in five ethnic groups: the Multiethnic Cohort. <i>British Journal of Nutrition</i> , 2017, 118, 312-320.	1.2	55
202	A promoter polymorphism in the CASP8 gene is not associated with cancer risk. <i>Nature Genetics</i> , 2008, 40, 259-260.	9.4	54
203	Mitochondrial DNA G10398A variant is not associated with breast cancer in African-American women. <i>Cancer Genetics and Cytogenetics</i> , 2008, 181, 16-19.	1.0	54
204	Meta-analysis of loci associated with age at natural menopause in African-American women. <i>Human Molecular Genetics</i> , 2014, 23, 3327-3342.	1.4	54
205	Prediction of individual genetic risk to prostate cancer using a polygenic score. <i>Prostate</i> , 2015, 75, 1467-1474.	1.2	54
206	A Haplotype-Based Case-Control Study of BRCA1 and Sporadic Breast Cancer Risk. <i>Cancer Research</i> , 2005, 65, 7516-7522.	0.4	53
207	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2014, 23, 6096-6111.	1.4	53
208	A Polygenic Risk Score for Breast Cancer in US Latinas and Latin American Women. <i>Journal of the National Cancer Institute</i> , 2020, 112, 590-598.	3.0	53
209	Genome-wide association study of age at menarche in African-American women. <i>Human Molecular Genetics</i> , 2013, 22, 3329-3346.	1.4	52
210	Racial/ethnic differences in lifestyle-related factors and prostate cancer risk: the Multiethnic Cohort Study. <i>Cancer Causes and Control</i> , 2015, 26, 1507-1515.	0.8	52
211	A Loss-of-Function Splice Acceptor Variant in <i>IGF2</i> Is Protective for Type 2 Diabetes. <i>Diabetes</i> , 2017, 66, 2903-2914.	0.3	52
212	Genome-wide association study of germline variants and breast cancer-specific mortality. <i>British Journal of Cancer</i> , 2019, 120, 647-657.	2.9	52
213	Common variation in BRCA2 and breast cancer risk: a haplotype-based analysis in the Multiethnic Cohort. <i>Human Molecular Genetics</i> , 2004, 13, 2431-2441.	1.4	51
214	Common Genetic Variants in Prostate Cancer Risk Predictionâ€”Results from the NCI Breast and Prostate Cancer Cohort Consortium (BPC3). <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 437-444.	1.1	51
215	Evaluating Genetic Risk for Prostate Cancer among Japanese and Latinos. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 2048-2058.	1.1	51
216	Large-scale association analysis in Asians identifies new susceptibility loci for prostate cancer. <i>Nature Communications</i> , 2015, 6, 8469.	5.8	51

#	ARTICLE	IF	CITATIONS
217	Fine-scale mapping of 8q24 locus identifies multiple independent risk variants for breast cancer. <i>International Journal of Cancer</i> , 2016, 139, 1303-1317.	2.3	51
218	Pancreatic Cancer Following Incident Diabetes in African Americans and Latinos: The Multiethnic Cohort. <i>Journal of the National Cancer Institute</i> , 2019, 111, 27-33.	3.0	51
219	Contribution of Germline Predisposition Gene Mutations to Breast Cancer Risk in African American Women. <i>Journal of the National Cancer Institute</i> , 2020, 112, 1213-1221.	3.0	51
220	Genetic Determinants of Circulating Insulin-Like Growth Factor (IGF)-I, IGF Binding Protein (BP)-1, and IGFBP-3 Levels in a Multiethnic Population. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2007, 92, 3660-3666.	1.8	50
221	Integration of multiethnic fine-mapping and genomic annotation to prioritize candidate functional SNPs at prostate cancer susceptibility regions. <i>Human Molecular Genetics</i> , 2015, 24, 5603-5618.	1.4	50
222	Genome-wide association studies in women of African ancestry identified 3q26.21 as a novel susceptibility locus for oestrogen receptor negative breast cancer. <i>Human Molecular Genetics</i> , 2016, 25, ddw305.	1.4	50
223	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. <i>Nature Communications</i> , 2016, 7, 10979.	5.8	50
224	Consistent Directions of Effect for Established Type 2 Diabetes Risk Variants Across Populations. <i>Diabetes</i> , 2012, 61, 1642-1647.	0.3	49
225	MicroRNA Related Polymorphisms and Breast Cancer Risk. <i>PLoS ONE</i> , 2014, 9, e109973.	1.1	49
226	Genome-wide association study confirms lung cancer susceptibility loci on chromosomes 5p15 and 15q25 in an African-American population. <i>Lung Cancer</i> , 2016, 98, 33-42.	0.9	49
227	12 new susceptibility loci for prostate cancer identified by genome-wide association study in Japanese population. <i>Nature Communications</i> , 2019, 10, 4422.	5.8	49
228	Determinants of penetrance and variable expressivity in monogenic metabolic conditions across 77,184 exomes. <i>Nature Communications</i> , 2021, 12, 3505.	5.8	49
229	A Large Study of Androgen Receptor Germline Variants and Their Relation to Sex Hormone Levels and Prostate Cancer Risk. Results from the National Cancer Institute Breast and Prostate Cancer Cohort Consortium. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2010, 95, E121-E127.	1.8	48
230	Quantifying the Genetic Correlation between Multiple Cancer Types. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1427-1435.	1.1	48
231	Race, ethnicity, community-level socioeconomic factors, and risk of COVID-19 in the United States and the United Kingdom. <i>EClinicalMedicine</i> , 2021, 38, 101029.	3.2	48
232	Multiple Polymorphisms in the Renin-Angiotensin-Aldosterone System (ACE, CYP11B2, AGTR1) and Their Contribution to Hypertension in African Americans and Latinos in the Multiethnic Cohort. <i>American Journal of the Medical Sciences</i> , 2004, 328, 266-273.	0.4	47
233	Haplotype-Based Association Studies of IGFBP1 and IGFBP3 with Prostate and Breast Cancer Risk: The Multiethnic Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 1993-1997.	1.1	47
234	A comprehensive analysis of common IGF1, IGFBP1 and IGFBP3 genetic variation with prospective IGF-I and IGFBP-3 blood levels and prostate cancer risk among Caucasians. <i>Human Molecular Genetics</i> , 2010, 19, 3089-3101.	1.4	47

#	ARTICLE	IF	CITATIONS
235	Rapid Assessment of Genetic Ancestry in Populations of Unknown Origin by Genome-Wide Genotyping of Pooled Samples. <i>PLoS Genetics</i> , 2010, 6, e1000866.	1.5	47
236	The Contribution of Common Genetic Variation to Nicotine and Cotinine Glucuronidation in Multiple Ethnic/Racial Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 119-127.	1.1	47
237	Risk of Breast Cancer Among Carriers of Pathogenic Variants in Breast Cancer Predisposition Genes Varies by Polygenic Risk Score. <i>Journal of Clinical Oncology</i> , 2021, 39, 2564-2573.	0.8	47
238	Established Risk Factors Account for Most of the Racial Differences in Cardiovascular Disease Mortality. <i>PLoS ONE</i> , 2007, 2, e377.	1.1	47
239	An Admixture Scan in 1,484 African American Women with Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 3110-3117.	1.1	46
240	Genetic variation in DNA repair pathway genes and premenopausal breast cancer risk. <i>Breast Cancer Research and Treatment</i> , 2009, 115, 613-622.	1.1	46
241	Comprehensive analysis of common genetic variation in 61 genes related to steroid hormone and insulin-like growth factor-I metabolism and breast cancer risk in the NCI breast and prostate cancer cohort consortium. <i>Human Molecular Genetics</i> , 2010, 19, 3873-3884.	1.4	45
242	Replication and validation of genetic polymorphisms associated with survival after allogeneic blood or marrow transplant. <i>Blood</i> , 2017, 130, 1585-1596.	0.6	45
243	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017, 46, 1814-1822.	0.9	45
244	Combined Associations of a Polygenic Risk Score and Classical Risk Factors With Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2021, 113, 329-337.	3.0	45
245	Mapping adipose and muscle tissue expression quantitative trait loci in African Americans to identify genes for type 2 diabetes and obesity. <i>Human Genetics</i> , 2016, 135, 869-880.	1.8	44
246	A functionally significant SNP in TP53 and breast cancer risk in African-American women. <i>Npj Breast Cancer</i> , 2017, 3, 5.	2.3	44
247	Ethnic differences in ovulatory function in nulliparous women. <i>British Journal of Cancer</i> , 2002, 86, 367-371.	2.9	43
248	Self-reported ethnicity, genetic structure and the impact of population stratification in a multiethnic study. <i>Human Genetics</i> , 2010, 128, 165-177.	1.8	43
249	Pooled Analysis of Phosphatidylinositol 3-Kinase Pathway Variants and Risk of Prostate Cancer. <i>Cancer Research</i> , 2010, 70, 2389-2396.	0.4	43
250	Type 2 diabetes risk variants and colorectal cancer risk: the Multiethnic Cohort and PAGE studies. <i>Gut</i> , 2011, 60, 1703-1711.	6.1	43
251	Fine mapping of a region of chromosome 11q13 reveals multiple independent loci associated with risk of prostate cancer. <i>Human Molecular Genetics</i> , 2011, 20, 2869-2878.	1.4	43
252	Genetic predisposition to ductal carcinoma in situ of the breast. <i>Breast Cancer Research</i> , 2016, 18, 22.	2.2	43

#	ARTICLE	IF	CITATIONS
253	Telomere structure and maintenance gene variants and risk of five cancer types. <i>International Journal of Cancer</i> , 2016, 139, 2655-2670.	2.3	43
254	Germline variation at 8q24 and prostate cancer risk in men of European ancestry. <i>Nature Communications</i> , 2018, 9, 4616.	5.8	43
255	Pathogenic and likely pathogenic variants in <i>PALB2</i> , <i>CHEK2</i> , and other known breast cancer susceptibility genes among 1054 BRCA-negative Hispanics with breast cancer. <i>Cancer</i> , 2019, 125, 2829-2836.	2.0	43
256	Identification of Novel Susceptibility Loci and Genes for Prostate Cancer Risk: A Transcriptome-Wide Association Study in Over 140,000 European Descendants. <i>Cancer Research</i> , 2019, 79, 3192-3204.	0.4	43
257	Comprehensive association testing of common genetic variation in DNA repair pathway genes in relationship with breast cancer risk in multiple populations. <i>Human Molecular Genetics</i> , 2008, 17, 825-834.	1.4	42
258	Levels of Beta-Microseminoprotein in Blood and Risk of Prostate Cancer in Multiple Populations. <i>Journal of the National Cancer Institute</i> , 2013, 105, 237-243.	3.0	42
259	A comprehensive examination of breast cancer risk loci in African American women. <i>Human Molecular Genetics</i> , 2014, 23, 5518-5526.	1.4	42
260	Genome-wide association study of endometrial cancer in E2C2. <i>Human Genetics</i> , 2014, 133, 211-224.	1.8	42
261	Pubertal development and prostate cancer risk: Mendelian randomization study in a population-based cohort. <i>BMC Medicine</i> , 2016, 14, 66.	2.3	42
262	Type II Diabetes, Obesity, and Breast Cancer Risk: The Multiethnic Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 854-861.	1.1	41
263	Evaluating Polygenic Risk Scores for Breast Cancer in Women of African Ancestry. <i>Journal of the National Cancer Institute</i> , 2021, 113, 1168-1176.	3.0	41
264	Comprehensive Functional Annotation of Seventy-One Breast Cancer Risk Loci. <i>PLoS ONE</i> , 2013, 8, e63925.	1.1	41
265	A High-Density Admixture Scan in 1,670 African Americans with Hypertension. <i>PLoS Genetics</i> , 2007, 3, e196.	1.5	40
266	A comprehensive analysis of common genetic variation in prolactin (PRL) and PRL receptor (PRLR) genes in relation to plasma prolactin levels and breast cancer risk: the Multiethnic Cohort. <i>BMC Medical Genetics</i> , 2007, 8, 72.	2.1	40
267	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , 2015, 24, 2966-2984.	1.4	40
268	Rare germline variants in DNA repair genes and the angiogenesis pathway predispose prostate cancer patients to develop metastatic disease. <i>British Journal of Cancer</i> , 2018, 119, 96-104.	2.9	40
269	Germline Sequencing DNA Repair Genes in 5545 Men With Aggressive and Nonaggressive Prostate Cancer. <i>Journal of the National Cancer Institute</i> , 2021, 113, 616-625.	3.0	40
270	Genetic Predisposition to In Situ and Invasive Lobular Carcinoma of the Breast. <i>PLoS Genetics</i> , 2014, 10, e1004285.	1.5	39

#	ARTICLE	IF	CITATIONS
271	Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. <i>American Journal of Human Genetics</i> , 2020, 107, 837-848.	2.6	39
272	Evaluation of a Multiethnic Polygenic Risk Score Model for Prostate Cancer. <i>Journal of the National Cancer Institute</i> , 2022, 114, 771-774.	3.0	39
273	Combined Effect of a Polygenic Risk Score and Rare Genetic Variants on Prostate Cancer Risk. <i>European Urology</i> , 2021, 80, 134-138.	0.9	39
274	Genetic Variation and Reproductive Timing: African American Women from the Population Architecture Using Genomics and Epidemiology (PAGE) Study. <i>PLoS ONE</i> , 2013, 8, e55258.	1.1	39
275	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. <i>Human Molecular Genetics</i> , 2015, 24, 285-298.	1.4	38
276	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. <i>Human Molecular Genetics</i> , 2016, 25, 1203-1214.	1.4	38
277	The Dietary Inflammatory Index and All-Cause, Cardiovascular Disease, and Cancer Mortality in the Multiethnic Cohort Study. <i>Nutrients</i> , 2018, 10, 1844.	1.7	38
278	Tobacco biomarkers and genetic/epigenetic analysis to investigate ethnic/racial differences in lung cancer risk among smokers. <i>Npj Precision Oncology</i> , 2018, 2, 17.	2.3	38
279	Quantitative trait loci predicting circulating sex steroid hormones in men from the NCI-Breast and Prostate Cancer Cohort Consortium (BPC3). <i>Human Molecular Genetics</i> , 2009, 18, 3749-3757.	1.4	37
280	Post-GWAS gene-environment interplay in breast cancer: results from the Breast and Prostate Cancer Cohort Consortium and a meta-analysis on 79 000 women. <i>Human Molecular Genetics</i> , 2014, 23, 5260-5270.	1.4	37
281	Polymorphisms in a Putative Enhancer at the 10q21.2 Breast Cancer Risk Locus Regulate NRBF2 Expression. <i>American Journal of Human Genetics</i> , 2015, 97, 22-34.	2.6	37
282	The genetic interplay between body mass index, breast size and breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2019, 48, 781-794.	0.9	37
283	Analysis of case-control association studies with known risk variants. <i>Bioinformatics</i> , 2012, 28, 1729-1737.	1.8	36
284	Genetic Variation in the Vitamin D Pathway in Relation to Risk of Prostate Cancer—Results from the Breast and Prostate Cancer Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 688-696.	1.1	36
285	Fine-mapping of genome-wide association study-identified risk loci for colorectal cancer in African Americans. <i>Human Molecular Genetics</i> , 2013, 22, 5048-5055.	1.4	36
286	Additive Interactions Between Susceptibility Single-Nucleotide Polymorphisms Identified in Genome-Wide Association Studies and Breast Cancer Risk Factors in the Breast and Prostate Cancer Cohort Consortium. <i>American Journal of Epidemiology</i> , 2014, 180, 1018-1027.	1.6	36
287	A tetranucleotide repeat polymorphism in CYP19 and breast cancer risk. <i>International Journal of Cancer</i> , 2000, 87, 204-210.	2.3	36
288	GWAS identifies a common breast cancer risk allele among BRCA1 carriers. <i>Nature Genetics</i> , 2010, 42, 819-820.	9.4	35

#	ARTICLE	IF	CITATIONS
289	Pleiotropic effects of genetic risk variants for other cancers on colorectal cancer risk: PAGE, GECCO and CCFR consortia. <i>Gut</i> , 2014, 63, 800-807.	6.1	35
290	Pleiotropic Associations of Risk Variants Identified for Other Cancers With Lung Cancer Risk: The PAGE and TRICL Consortia. <i>Journal of the National Cancer Institute</i> , 2014, 106, dju061.	3.0	35
291	Interethnic differences in pancreatic cancer incidence and risk factors: The Multiethnic Cohort. <i>Cancer Medicine</i> , 2019, 8, 3592-3603.	1.3	35
292	Alcohol Intake and Colorectal Cancer Risk in the Multiethnic Cohort Study. <i>American Journal of Epidemiology</i> , 2019, 188, 67-76.	1.6	35
293	Mendelian randomization analyses suggest a role for cholesterol in the development of endometrial cancer. <i>International Journal of Cancer</i> , 2021, 148, 307-319.	2.3	35
294	HNF1B and Endometrial Cancer Risk: Results from the PAGE study. <i>PLoS ONE</i> , 2012, 7, e30390.	1.1	34
295	Genetic Ancestry and Risk of Mortality among U.S. Latinas with Breast Cancer. <i>Cancer Research</i> , 2013, 73, 7243-7253.	0.4	34
296	Fine-Mapping the HOXB Region Detects Common Variants Tagging a Rare Coding Allele: Evidence for Synthetic Association in Prostate Cancer. <i>PLoS Genetics</i> , 2014, 10, e1004129.	1.5	34
297	Genome-wide association study of colorectal cancer in Hispanics. <i>Carcinogenesis</i> , 2016, 37, 547-556.	1.3	34
298	Circulating sex hormones in relation to anthropometric, sociodemographic and behavioural factors in an international dataset of 12,300 men. <i>PLoS ONE</i> , 2017, 12, e0187741.	1.1	34
299	The COronavirus Pandemic Epidemiology (COPE) Consortium: A Call to Action. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1283-1289.	1.1	34
300	Genome-wide association meta-analysis identifies GP2 gene risk variants for pancreatic cancer. <i>Nature Communications</i> , 2020, 11, 3175.	5.8	34
301	<i>CYP19A1</i> Genetic Variation in Relation to Prostate Cancer Risk and Circulating Sex Hormone Concentrations in Men from the Breast and Prostate Cancer Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 2734-2744.	1.1	33
302	Association of breast cancer risk loci with breast cancer survival. <i>International Journal of Cancer</i> , 2015, 137, 2837-2845.	2.3	33
303	An intergenic risk locus containing an enhancer deletion in 2q35 modulates breast cancer risk by deregulating IGFBP5 expression. <i>Human Molecular Genetics</i> , 2016, 25, 3863-3876.	1.4	33
304	A rare variant of African ancestry activates 8q24 lncRNA hub by modulating cancer associated enhancer. <i>Nature Communications</i> , 2020, 11, 3598.	5.8	33
305	Urinary phthalate exposures and risk of breast cancer: the Multiethnic Cohort study. <i>Breast Cancer Research</i> , 2021, 23, 44.	2.2	33
306	Diabetes and urothelial cancer risk: The Multiethnic Cohort Study. <i>Cancer Epidemiology</i> , 2011, 35, 551-554.	0.8	32

#	ARTICLE	IF	CITATIONS
307	A large-scale assessment of two-way SNP interactions in breast cancer susceptibility using 46 450 cases and 42 461 controls from the breast cancer association consortium. <i>Human Molecular Genetics</i> , 2014, 23, 1934-1946.	1.4	32
308	Genetic variants in microRNA and microRNA biogenesis pathway genes and breast cancer risk among women of African ancestry. <i>Human Genetics</i> , 2016, 135, 1145-1159.	1.8	32
309	Replication of associations between genetic polymorphisms and chronic graft-versus-host disease. <i>Blood</i> , 2016, 128, 2450-2456.	0.6	32
310	Identification of novel common breast cancer risk variants at the 6q25 locus among Latinas. <i>Breast Cancer Research</i> , 2019, 21, 3.	2.2	32
311	A Germline Variant at 8q24 Contributes to Familial Clustering of Prostate Cancer in Men of African Ancestry. <i>European Urology</i> , 2020, 78, 316-320.	0.9	32
312	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	0.6	32
313	Deconstructing, Addressing, and Eliminating Racial and Ethnic Inequities in Prostate Cancer Care. <i>European Urology</i> , 2022, 82, 341-351.	0.9	32
314	N-Acetyltransferase 2 Polymorphisms, Tobacco Smoking, and Breast Cancer Risk in the Breast and Prostate Cancer Cohort Consortium. <i>American Journal of Epidemiology</i> , 2011, 174, 1316-1322.	1.6	31
315	Susceptibility variants for obesity and colorectal cancer risk: The multiethnic cohort and PAGE studies. <i>International Journal of Cancer</i> , 2012, 131, E1038-43.	2.3	31
316	Multiancestral Analysis of Inflammation-Related Genetic Variants and C-Reactive Protein in the Population Architecture Using Genomics and Epidemiology Study. <i>Circulation: Cardiovascular Genetics</i> , 2014, 7, 178-188.	5.1	31
317	Association of breast cancer risk with genetic variants showing differential allelic expression: Identification of a novel breast cancer susceptibility locus at 4q21. <i>Oncotarget</i> , 2016, 7, 80140-80163.	0.8	31
318	Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. <i>Breast Cancer Research</i> , 2016, 18, 64.	2.2	31
319	Trans-ethnic fine-mapping of genetic loci for body mass index in the diverse ancestral populations of the Population Architecture using Genomics and Epidemiology (PAGE) Study reveals evidence for multiple signals at established loci. <i>Human Genetics</i> , 2017, 136, 771-800.	1.8	31
320	Genetic risk of prostate cancer in Ugandan men. <i>Prostate</i> , 2018, 78, 370-376.	1.2	31
321	Joint association of mammographic density adjusted for age and body mass index and polygenic risk score with breast cancer risk. <i>Breast Cancer Research</i> , 2019, 21, 68.	2.2	31
322	Protein-altering germline mutations implicate novel genes related to lung cancer development. <i>Nature Communications</i> , 2020, 11, 2220.	5.8	31
323	Polymorphisms in genes involved in estrogen and progesterone metabolism and mammographic density changes in women randomized to postmenopausal hormone therapy: results from a pilot study. <i>Breast Cancer Research</i> , 2005, 7, R336-44.	2.2	30
324	Hormone-related pathways and risk of breast cancer subtypes in African American women. <i>Breast Cancer Research and Treatment</i> , 2015, 154, 145-154.	1.1	30

#	ARTICLE	IF	CITATIONS
325	Smoking and breast cancer risk by race/ethnicity and oestrogen and progesterone receptor status: the Multiethnic Cohort (MEC) study. <i>International Journal of Epidemiology</i> , 2019, 48, 501-511.	0.9	30
326	A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. <i>Nature Communications</i> , 2020, 11, 312.	5.8	30
327	Pathogenic Variants in Cancer Predisposition Genes and Prostate Cancer Risk in Men of African Ancestry. <i>JCO Precision Oncology</i> , 2020, 4, 32-43.	1.5	30
328	Myth: Blood transfusion is effective for sickle cell anemia-associated priapism. <i>Canadian Journal of Emergency Medicine</i> , 2006, 8, 119-122.	0.5	29
329	The Influence of Obesity-Related Single Nucleotide Polymorphisms on BMI Across the Life Course: The PAGE Study. <i>Diabetes</i> , 2013, 62, 1763-1767.	0.3	29
330	Fine-mapping of lipid regions in global populations discovers ethnic-specific signals and refines previously identified lipid loci. <i>Human Molecular Genetics</i> , 2016, 25, 5500-5512.	1.4	29
331	Genetic susceptibility to radiation-induced breast cancer after Hodgkin lymphoma. <i>Blood</i> , 2019, 133, 1130-1139.	0.6	29
332	Exploring genetic susceptibility to cancer in diverse populations. <i>Current Opinion in Genetics and Development</i> , 2010, 20, 330-335.	1.5	28
333	Large-scale fine mapping of the HNF1B locus and prostate cancer risk. <i>Human Molecular Genetics</i> , 2011, 20, 3322-3329.	1.4	28
334	Insulin-like growth factor pathway genes and blood concentrations, dietary protein and risk of prostate cancer in the NCI Breast and Prostate Cancer Cohort Consortium (BPC3). <i>International Journal of Cancer</i> , 2013, 133, 495-504.	2.3	28
335	Alcohol consumption and prostate cancer incidence and progression: A Mendelian randomisation study. <i>International Journal of Cancer</i> , 2017, 140, 75-85.	2.3	28
336	Prospective Study of Coffee Consumption and Cancer Incidence in Non-White Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 928-935.	1.1	28
337	The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. <i>Npj Breast Cancer</i> , 2019, 5, 38.	2.3	28
338	An integrative multi-omics analysis to identify candidate DNA methylation biomarkers related to prostate cancer risk. <i>Nature Communications</i> , 2020, 11, 3905.	5.8	28
339	Common Genetic Variation at PTEN and Risk of Sporadic Breast and Prostate Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 1021-1025.	1.1	27
340	Association between breast cancer susceptibility loci and mammographic density: the Multiethnic Cohort. <i>Breast Cancer Research</i> , 2009, 11, R10.	2.2	27
341	Association of Genes, Pathways, and Haplogroups of the Mitochondrial Genome with the Risk of Colorectal Cancer: The Multiethnic Cohort. <i>PLoS ONE</i> , 2015, 10, e0136796.	1.1	27
342	Genome-Wide Association Study of Prostate Cancer-Specific Survival. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1796-1800.	1.1	27

#	ARTICLE	IF	CITATIONS
343	The causal relevance of body mass index in different histological types of lung cancer: A Mendelian randomization study. <i>Scientific Reports</i> , 2016, 6, 31121.	1.6	27
344	Low diet quality and the risk of stroke mortality: the multiethnic cohort study. <i>European Journal of Clinical Nutrition</i> , 2018, 72, 1035-1045.	1.3	27
345	Population-specific reference panels are crucial for genetic analyses: an example of the CREBRF locus in Native Hawaiians. <i>Human Molecular Genetics</i> , 2020, 29, 2275-2284.	1.4	27
346	Variation in genetic admixture and population structure among Latinos: the Los Angeles Latino eye study (LALES). <i>BMC Genetics</i> , 2009, 10, 71.	2.7	26
347	Refining the Prostate Cancer Genetic Association within the <i>JAZF1</i> Gene on Chromosome 7p15.2. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 1349-1355.	1.1	26
348	Common germline polymorphisms associated with breast cancer-specific survival. <i>Breast Cancer Research</i> , 2015, 17, 58.	2.2	26
349	Assessing the role of insulin-like growth factors and binding proteins in prostate cancer using Mendelian randomization: Genetic variants as instruments for circulating levels. <i>International Journal of Cancer</i> , 2016, 139, 1520-1533.	2.3	26
350	Whole-exome sequencing of over 4100 men of African ancestry and prostate cancer risk. <i>Human Molecular Genetics</i> , 2016, 25, 371-381.	1.4	26
351	Novel colon cancer susceptibility variants identified from a genome-wide association study in African Americans. <i>International Journal of Cancer</i> , 2017, 140, 2728-2733.	2.3	26
352	Characterizing the neighborhood obesogenic environment in the Multiethnic Cohort: a multi-level infrastructure for cancer health disparities research. <i>Cancer Causes and Control</i> , 2018, 29, 167-183.	0.8	26
353	Sleep duration and incidence of type 2 diabetes: the Multiethnic Cohort. <i>Sleep Health</i> , 2018, 4, 27-32.	1.3	26
354	Smoking-Related Risks of Colorectal Cancer by Anatomical Subsite and Sex. <i>American Journal of Epidemiology</i> , 2020, 189, 543-553.	1.6	26
355	RAD51B in Familial Breast Cancer. <i>PLoS ONE</i> , 2016, 11, e0153788.	1.1	26
356	Genetic risk variants associated with in situ breast cancer. <i>Breast Cancer Research</i> , 2015, 17, 82.	2.2	25
357	The genetic underpinnings of variation in ages at menarche and natural menopause among women from the multi-ethnic Population Architecture using Genomics and Epidemiology (PAGE) Study: A trans-ethnic meta-analysis. <i>PLoS ONE</i> , 2018, 13, e0200486.	1.1	25
358	Prediagnostic circulating markers of inflammation and risk of oesophageal adenocarcinoma: a study within the National Cancer Institute Cohort Consortium. <i>Gut</i> , 2019, 68, 960-968.	6.1	25
359	A Collaborative Analysis of Individual Participant Data from 19 Prospective Studies Assesses Circulating Vitamin D and Prostate Cancer Risk. <i>Cancer Research</i> , 2019, 79, 274-285.	0.4	25
360	Screening and association testing of common coding variation in steroid hormone receptor co-activator and co-repressor genes in relation to breast cancer risk: the Multiethnic Cohort. <i>BMC Cancer</i> , 2009, 9, 43.	1.1	24

#	ARTICLE	IF	CITATIONS
361	Characterization of the colorectal cancer-associated enhancer MYC-335 at 8q24: the role of rs67491583. <i>Cancer Genetics</i> , 2012, 205, 25-33.	0.2	24
362	Genetic variants associated with fasting glucose and insulin concentrations in an ethnically diverse population: results from the Population Architecture using Genomics and Epidemiology (PAGE) study. <i>BMC Medical Genetics</i> , 2013, 14, 98.	2.1	24
363	The 19q12 Bladder Cancer GWAS Signal: Association with Cyclin E Function and Aggressive Disease. <i>Cancer Research</i> , 2014, 74, 5808-5818.	0.4	24
364	Fine-Scale Mapping of the 4q24 Locus Identifies Two Independent Loci Associated with Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1680-1691.	1.1	24
365	Characterizing Genetic Susceptibility to Breast Cancer in Women of African Ancestry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1016-1026.	1.1	24
366	Prediction and clinical utility of a contralateral breast cancer risk model. <i>Breast Cancer Research</i> , 2019, 21, 144.	2.2	24
367	A genome-wide association study of prostate cancer in Latinos. <i>International Journal of Cancer</i> , 2020, 146, 1819-1826.	2.3	24
368	Diet Quality and Risk of Lung Cancer in the Multiethnic Cohort Study. <i>Nutrients</i> , 2021, 13, 1614.	1.7	24
369	Cross-ancestry GWAS meta-analysis identifies six breast cancer loci in African and European ancestry women. <i>Nature Communications</i> , 2021, 12, 4198.	5.8	24
370	Replication of Five Prostate Cancer Loci Identified in an Asian Population—Results from the NCI Breast and Prostate Cancer Cohort Consortium (BPC3). <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012, 21, 212-216.	1.1	23
371	Association of Cancer Susceptibility Variants with Risk of Multiple Primary Cancers: The Population Architecture using Genomics and Epidemiology Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 2568-2578.	1.1	23
372	Metabolites of the Polycyclic Aromatic Hydrocarbon Phenanthrene in the Urine of Cigarette Smokers from Five Ethnic Groups with Differing Risks for Lung Cancer. <i>PLoS ONE</i> , 2016, 11, e0156203.	1.1	23
373	Polyunsaturated fatty acids and prostate cancer risk: a Mendelian randomisation analysis from the PRACTICAL consortium. <i>British Journal of Cancer</i> , 2016, 115, 624-631.	2.9	23
374	Dietary Patterns Derived by Reduced Rank Regression Are Inversely Associated with Type 2 Diabetes Risk across 5 Ethnic Groups in the Multiethnic Cohort. <i>Current Developments in Nutrition</i> , 2017, 1, e000620.	0.1	23
375	High-Quality Diets Are Associated With Reduced Risk of Hepatocellular Carcinoma and Chronic Liver Disease: The Multiethnic Cohort. <i>Hepatology Communications</i> , 2019, 3, 437-447.	2.0	23
376	Genetic discovery and risk characterization in type 2 diabetes across diverse populations. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100029.	1.0	23
377	Polygenic risk modeling for prediction of epithelial ovarian cancer risk. <i>European Journal of Human Genetics</i> , 2022, 30, 349-362.	1.4	23
378	A Healthy Lifestyle in Men at Increased Genetic Risk for Prostate Cancer. <i>European Urology</i> , 2023, 83, 343-351.	0.9	23

#	ARTICLE	IF	CITATIONS
379	An exome-wide analysis of low frequency and rare variants in relation to risk of breast cancer in African American Women: the AMBER Consortium. <i>Carcinogenesis</i> , 2016, 37, 870-877.	1.3	22
380	The Four-Kallikrein Panel Is Effective in Identifying Aggressive Prostate Cancer in a Multiethnic Population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1381-1388.	1.1	22
381	Association of Genetic Risk Score With NAFLD in An Ethnically Diverse Cohort. <i>Hepatology Communications</i> , 2021, 5, 1689-1703.	2.0	22
382	Germline Pathogenic Variants in Cancer Predisposition Genes Among Women With Invasive Lobular Carcinoma of the Breast. <i>Journal of Clinical Oncology</i> , 2021, 39, 3918-3926.	0.8	22
383	A Rare Germline HOXB13 Variant Contributes to Risk of Prostate Cancer in Men of African Ancestry. <i>European Urology</i> , 2022, 81, 458-462.	0.9	22
384	Polymorphic repeat in AIB1 does not alter breast cancer risk. <i>Breast Cancer Research</i> , 2000, 2, 378-85.	2.2	21
385	Use of weighted reference panels based on empirical estimates of ancestry for capturing untyped variation. <i>Human Genetics</i> , 2009, 125, 295-303.	1.8	21
386	A Genome-wide Pleiotropy Scan for Prostate Cancer Risk. <i>European Urology</i> , 2015, 67, 649-657.	0.9	21
387	Association of genetic susceptibility variants for type 2 diabetes with breast cancer risk in women of European ancestry. <i>Cancer Causes and Control</i> , 2016, 27, 679-693.	0.8	21
388	Genetic variations in vitamin D-related pathways and breast cancer risk in African American women in the AMBER consortium. <i>International Journal of Cancer</i> , 2016, 138, 2118-2126.	2.3	21
389	Exome chip analyses identify genes affecting mortality after HLA-matched unrelated-donor blood and marrow transplantation. <i>Blood</i> , 2018, 131, 2490-2499.	0.6	21
390	Antibody Responses to Streptococcus Gallolyticus Subspecies Gallolyticus Proteins in a Large Prospective Colorectal Cancer Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 1186-1194.	1.1	21
391	Agnostic Pathway/Gene Set Analysis of Genome-Wide Association Data Identifies Associations for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , 2019, 111, 557-567.	3.0	21
392	Circulating Metabolic Biomarkers of Screen-Detected Prostate Cancer in the ProtecT Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 208-216.	1.1	21
393	Circulating markers of cellular immune activation in prediagnostic blood sample and lung cancer risk in the Lung Cancer Cohort Consortium (LC3). <i>International Journal of Cancer</i> , 2020, 146, 2394-2405.	2.3	21
394	Genome-Wide Association Study of Liver Fat: The Multiethnic Cohort Adiposity Phenotype Study. <i>Hepatology Communications</i> , 2020, 4, 1112-1123.	2.0	21
395	Risk of breast cancer and prediagnostic urinary excretion of bisphenol A, triclosan and parabens: The Multiethnic Cohort Study. <i>International Journal of Cancer</i> , 2021, 149, 1426-1434.	2.3	21
396	Risk of Late-Onset Breast Cancer in Genetically Predisposed Women. <i>Journal of Clinical Oncology</i> , 2021, 39, 3430-3440.	0.8	21

#	ARTICLE	IF	CITATIONS
397	Comparison of the Prevalence of Pathogenic Variants in Cancer Susceptibility Genes in Black Women and Non-Hispanic White Women With Breast Cancer in the United States. <i>JAMA Oncology</i> , 2021, 7, 1045.	3.4	21
398	Genetic variation in the progesterone receptor gene and risk of endometrial cancer: a haplotype-based approach. <i>Carcinogenesis</i> , 2010, 31, 1392-1399.	1.3	20
399	8q24 risk alleles and prostate cancer in African-Barbadian men. <i>Prostate</i> , 2014, 74, 1579-1588.	1.2	20
400	SNP-SNP interaction analysis of NF- κ B signaling pathway on breast cancer survival. <i>Oncotarget</i> , 2015, 6, 37979-37994.	0.8	20
401	Vitamin D-Associated Genetic Variation and Risk of Breast Cancer in the Breast and Prostate Cancer Cohort Consortium (BPC3). <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 627-630.	1.1	20
402	Genetic variations in the Hippo signaling pathway and breast cancer risk in African American women in the AMBER Consortium. <i>Carcinogenesis</i> , 2016, 37, 951-956.	1.3	20
403	Transethnic insight into the genetics of glycaemic traits: fine-mapping results from the Population Architecture using Genomics and Epidemiology (PAGE) consortium. <i>Diabetologia</i> , 2017, 60, 2384-2398.	2.9	20
404	The impact of global and local Polynesian genetic ancestry on complex traits in Native Hawaiians. <i>PLoS Genetics</i> , 2021, 17, e1009273.	1.5	20
405	Benzene Uptake and Glutathione S-transferase T1 Status as Determinants of S-Phenylmercapturic Acid in Cigarette Smokers in the Multiethnic Cohort. <i>PLoS ONE</i> , 2016, 11, e0150641.	1.1	20
406	A Systematic Assessment of Common Genetic Variation in CYP11A and Risk of Breast Cancer. <i>Cancer Research</i> , 2006, 66, 12019-12025.	0.4	19
407	A comprehensive evaluation of interaction between genetic variants and use of menopausal hormone therapy on mammographic density. <i>Breast Cancer Research</i> , 2015, 17, 110.	2.2	19
408	Interaction between Red Meat Intake and NAT2 Genotype in Increasing the Risk of Colorectal Cancer in Japanese and African Americans. <i>PLoS ONE</i> , 2015, 10, e0144955.	1.1	19
409	Admixture Mapping of African-American Women in the AMBER Consortium Identifies New Loci for Breast Cancer and Estrogen-Receptor Subtypes. <i>Frontiers in Genetics</i> , 2016, 7, 170.	1.1	19
410	Fine scale mapping of the 17q22 breast cancer locus using dense SNPs, genotyped within the Collaborative Oncological Gene-Environment Study (COGs). <i>Scientific Reports</i> , 2016, 6, 32512.	1.6	19
411	Postmenopausal Female Hormone Use and Estrogen Receptor-Positive and -Negative Breast Cancer in African American Women. <i>Journal of the National Cancer Institute</i> , 2016, 108, djv361.	3.0	19
412	GWAS meta-analysis of 16 852 women identifies new susceptibility locus for endometrial cancer. <i>Human Molecular Genetics</i> , 2016, 25, ddw092.	1.4	19
413	Fine mapping of QT interval regions in global populations refines previously identified QT interval loci and identifies signals unique to African and Hispanic descent populations. <i>Heart Rhythm</i> , 2017, 14, 572-580.	0.3	19
414	The <i>BRCA2</i> c.68-7T>A variant is not pathogenic: A model for clinical calibration of spliceogenicity. <i>Human Mutation</i> , 2018, 39, 729-741.	1.1	19

#	ARTICLE	IF	CITATIONS
415	Type 2 diabetes and colorectal cancer survival: The multiethnic cohort. <i>International Journal of Cancer</i> , 2018, 143, 263-268.	2.3	19
416	A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. <i>Nature Communications</i> , 2021, 12, 1078.	5.8	19
417	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 623-642.	1.1	19
418	Population Genetic Structure and Origins of Native Hawaiians in the Multiethnic Cohort Study. <i>PLoS ONE</i> , 2012, 7, e47881.	1.1	19
419	A Variant in the Cytochrome P450 Oxidoreductase Gene Is Associated with Breast Cancer Risk in African Americans. <i>Cancer Research</i> , 2007, 67, 3565-3568.	0.4	18
420	Cross-cancer pleiotropic analysis of endometrial cancer: PAGE and E2C2 consortia. <i>Carcinogenesis</i> , 2014, 35, 2068-2073.	1.3	18
421	A Meta-analysis of Multiple Myeloma Risk Regions in African and European Ancestry Populations Identifies Putatively Functional Loci. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1609-1618.	1.1	18
422	No clinical utility of KRAS variant rs61764370 for ovarian or breast cancer. <i>Gynecologic Oncology</i> , 2016, 141, 386-401.	0.6	18
423	Type 2 diabetes as a predictor of survival among breast cancer patients: the multiethnic cohort. <i>Breast Cancer Research and Treatment</i> , 2019, 173, 637-645.	1.1	18
424	Identification of novel epithelial ovarian cancer loci in women of African ancestry. <i>International Journal of Cancer</i> , 2020, 146, 2987-2998.	2.3	18
425	Racial Differences in <i>Helicobacter pylori</i> CagA Sero-prevalence in a Consortium of Adult Cohorts in the United States. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2084-2092.	1.1	18
426	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. <i>American Journal of Human Genetics</i> , 2021, 108, 564-582.	2.6	18
427	Associations between Genetically Predicted Blood Protein Biomarkers and Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1501-1508.	1.1	18
428	No association between a single nucleotide polymorphism in CYP19 and breast cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2002, 11, 215-6.	1.1	18
429	Risk of Alzheimer's disease and related dementia by sex and race/ethnicity: The Multiethnic Cohort Study. <i>Alzheimer's and Dementia</i> , 2022, 18, 1625-1634.	0.4	18
430	Circulating free testosterone and risk of aggressive prostate cancer: Prospective and Mendelian randomisation analyses in international consortia. <i>International Journal of Cancer</i> , 2022, 151, 1033-1046.	2.3	18
431	Utilizing HapMap and Tagging SNPs. <i>Methods in Molecular Medicine</i> , 2008, 141, 37-54.	0.8	17
432	A Common Prostate Cancer Risk Variant 5â€² of <i>Microseminoprotein-Î² (MSMB)</i> Is a Strong Predictor of Circulating Î²-Microseminoprotein (MSP) Levels in Multiple Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010, 19, 2639-2646.	1.1	17

#	ARTICLE	IF	CITATIONS
433	A systematic approach to understand the functional consequences of non-protein coding risk regions. <i>Cell Cycle</i> , 2010, 9, 256-259.	1.3	17
434	Caution in generalizing known genetic risk markers for breast cancer across all ethnic/racial populations. <i>European Journal of Human Genetics</i> , 2011, 19, 243-245.	1.4	17
435	Identification of shared and unique susceptibility pathways among cancers of the lung, breast, and prostate from genome-wide association studies and tissue-specific protein interactions. <i>Human Molecular Genetics</i> , 2015, 24, 7406-7420.	1.4	17
436	Gene and pathway level analyses of germline DNA-repair gene variants and prostate cancer susceptibility using the iCOGS-genotyping array. <i>British Journal of Cancer</i> , 2016, 114, 945-952.	2.9	17
437	Investigating the possible causal role of coffee consumption with prostate cancer risk and progression using Mendelian randomization analysis. <i>International Journal of Cancer</i> , 2017, 140, 322-328.	2.3	17
438	Pleiotropy of Cancer Susceptibility Variants on the Risk of Non-Hodgkin Lymphoma: The PAGE Consortium. <i>PLoS ONE</i> , 2014, 9, e89791.	1.1	16
439	2q36.3 is associated with prognosis for oestrogen receptor-negative breast cancer patients treated with chemotherapy. <i>Nature Communications</i> , 2014, 5, 4051.	5.8	16
440	Reducing GWAS Complexity. <i>Cell Cycle</i> , 2016, 15, 22-24.	1.3	16
441	Genetic Variants in Immune-Related Pathways and Breast Cancer Risk in African American Women in the AMBER Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 321-330.	1.1	16
442	Discovery, fine-mapping, and conditional analyses of genetic variants associated with C-reactive protein in multiethnic populations using the MetaboChip in the Population Architecture using Genomics and Epidemiology (PAGE) study. <i>Human Molecular Genetics</i> , 2018, 27, 2940-2953.	1.4	16
443	Analysis of Over 140,000 European Descendants Identifies Genetically Predicted Blood Protein Biomarkers Associated with Prostate Cancer Risk. <i>Cancer Research</i> , 2019, 79, 4592-4598.	0.4	16
444	The CHEK2 Variant C.349A>G Is Associated with Prostate Cancer Risk and Carriers Share a Common Ancestor. <i>Cancers</i> , 2020, 12, 3254.	1.7	16
445	Additional SNPs improve risk stratification of a polygenic hazard score for prostate cancer. <i>Prostate Cancer and Prostatic Diseases</i> , 2021, 24, 532-541.	2.0	16
446	A meta-analysis of genome-wide association studies of multiple myeloma among men and women of African ancestry. <i>Blood Advances</i> , 2020, 4, 181-190.	2.5	16
447	Plasma Carotenoid- and Retinol-Weighted Multi-SNP Scores and Risk of Breast Cancer in the National Cancer Institute Breast and Prostate Cancer Cohort Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 927-936.	1.1	15
448	No Association between the Mitochondrial Genome and Prostate Cancer Risk: The Multiethnic Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1001-1003.	1.1	15
449	Genetic association with B-cell acute lymphoblastic leukemia in allogeneic transplant patients differs by age and sex. <i>Blood Advances</i> , 2017, 1, 1717-1728.	2.5	15
450	Circulating cotinine concentrations and lung cancer risk in the Lung Cancer Cohort Consortium (LC3). <i>International Journal of Epidemiology</i> , 2018, 47, 1760-1771.	0.9	15

#	ARTICLE	IF	CITATIONS
451	A survey of microRNA single nucleotide polymorphisms identifies novel breast cancer susceptibility loci in a case-control, population-based study of African-American women. <i>Breast Cancer Research</i> , 2018, 20, 45.	2.2	15
452	A phenome-wide association study (PheWAS) in the Population Architecture using Genomics and Epidemiology (PAGE) study reveals potential pleiotropy in African Americans. <i>PLoS ONE</i> , 2019, 14, e0226771.	1.1	15
453	Common variants in breast cancer risk loci predispose to distinct tumor subtypes. <i>Breast Cancer Research</i> , 2022, 24, 2.	2.2	15
454	Multi-ethnic GWAS and fine-mapping of glycaemic traits identify novel loci in the PAGE Study. <i>Diabetologia</i> , 2022, 65, 477-489.	2.9	15
455	Genome-wide and transcriptome-wide association studies of mammographic density phenotypes reveal novel loci. <i>Breast Cancer Research</i> , 2022, 24, 27.	2.2	15
456	Validation of a multi-ancestry polygenic risk score and age-specific risks of prostate cancer: A meta-analysis within diverse populations. <i>ELife</i> , 0, 11, .	2.8	15
457	ABO blood group alleles and prostate cancer risk: Results from the breast and prostate cancer cohort consortium (BPC3). <i>Prostate</i> , 2015, 75, 1677-1681.	1.2	14
458	Inherited variants in the inner centromere protein (INCENP) gene of the chromosomal passenger complex contribute to the susceptibility of ER-negative breast cancer. <i>Carcinogenesis</i> , 2015, 36, 256-271.	1.3	14
459	Physical Activity and Colorectal Cancer Risk by Sex, Race/Ethnicity, and Subsite: The Multiethnic Cohort Study. <i>Cancer Prevention Research</i> , 2019, 12, 315-326.	0.7	14
460	The associations of anthropometric, behavioural and sociodemographic factors with circulating concentrations of IGF1, IGF2, IGFBP1, IGFBP2 and IGFBP3 in a pooled analysis of 16,024 men from 22 studies. <i>International Journal of Cancer</i> , 2019, 145, 3244-3256.	2.3	14
461	Prediction of contralateral breast cancer: external validation of risk calculators in 20 international cohorts. <i>Breast Cancer Research and Treatment</i> , 2020, 181, 423-434.	1.1	14
462	Modifiable Factors and Incident Gout Across Ethnicity Within a Large Multiethnic Cohort of Older Adults. <i>Journal of Rheumatology</i> , 2022, 49, 504-512.	1.0	14
463	Prostate cancer risk stratification improvement across multiple ancestries with new polygenic hazard score. <i>Prostate Cancer and Prostatic Diseases</i> , 2022, 25, 755-761.	2.0	14
464	Joint Associations of Race, Ethnicity, and Socioeconomic Status With Mortality in the Multiethnic Cohort Study. <i>JAMA Network Open</i> , 2022, 5, e226370.	2.8	14
465	Distinct Reproductive Risk Profiles for Intrinsic-Like Breast Cancer Subtypes: Pooled Analysis of Population-Based Studies. <i>Journal of the National Cancer Institute</i> , 2022, 114, 1706-1719.	3.0	14
466	IGF1, IGFBP1, and IGFBP3 genes and mammographic density: The Multiethnic Cohort. <i>International Journal of Cancer</i> , 2010, 127, 1115-1123.	2.3	13
467	Body Mass Index Genetic Risk Score and Endometrial Cancer Risk. <i>PLoS ONE</i> , 2015, 10, e0143256.	1.1	13
468	Atopic Allergic Conditions and Colorectal Cancer Risk in the Multiethnic Cohort Study. <i>American Journal of Epidemiology</i> , 2015, 181, 889-897.	1.6	13

#	ARTICLE	IF	CITATIONS
469	Evaluation of 71 Coronary Artery Disease Risk Variants in a Multiethnic Cohort. <i>Frontiers in Cardiovascular Medicine</i> , 2018, 5, 19.	1.1	13
470	Techniques and Outcomes of MRI-TRUS Fusion Prostate Biopsy. <i>Current Urology Reports</i> , 2021, 22, 27.	1.0	13
471	Exome-Wide Association Study of Endometrial Cancer in a Multiethnic Population. <i>PLoS ONE</i> , 2014, 9, e97045.	1.1	12
472	Genetic variation in mitotic regulatory pathway genes is associated with breast tumor grade. <i>Human Molecular Genetics</i> , 2014, 23, 6034-6046.	1.4	12
473	Associations Between Genetic Ancestries and Nicotine Metabolism Biomarkers in the Multiethnic Cohort Study. <i>American Journal of Epidemiology</i> , 2015, 182, 945-951.	1.6	12
474	Fine-Mapping of the 1p11.2 Breast Cancer Susceptibility Locus. <i>PLoS ONE</i> , 2016, 11, e0160316.	1.1	12
475	A case-control analysis of smoking and breast cancer in African American women: findings from the AMBER Consortium. <i>Carcinogenesis</i> , 2016, 37, 607-615.	1.3	12
476	Gene-based analysis of the fibroblast growth factor receptor signaling pathway in relation to breast cancer in African American women: the AMBER consortium. <i>Breast Cancer Research and Treatment</i> , 2016, 155, 355-363.	1.1	12
477	Exploring Differences in the Aspirin-Colorectal Cancer Association by Sex and Race/Ethnicity: The Multiethnic Cohort Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 162-169.	1.1	12
478	Association between sleep duration and breast cancer incidence: The multiethnic cohort. <i>International Journal of Cancer</i> , 2020, 146, 664-670.	2.3	12
479	Diet quality and all-cause and cancer-specific mortality in cancer survivors and non-cancer individuals: the Multiethnic Cohort Study. <i>European Journal of Nutrition</i> , 2022, 61, 925-933.	1.8	12
480	No Association of Type 2 Diabetes Risk Variants and Prostate Cancer Risk: the Multiethnic Cohort and PAGE: Table 1.. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 1979-1981.	1.1	11
481	No Association of Risk Variants for Diabetes and Obesity with Breast Cancer: The Multiethnic Cohort and PAGE Studies: Table 1.. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 1039-1042.	1.1	11
482	SNP interaction pattern identifier (SIPI): an intensive search for SNP-SNP interaction patterns. <i>Bioinformatics</i> , 2017, 33, 822-833.	1.8	11
483	Replication and Genetic Risk Score Analysis for Pancreatic Cancer in a Diverse Multiethnic Population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2686-2692.	1.1	11
484	Polygenic risk scores for prediction of breast cancer risk in women of African ancestry: a cross-ancestry approach. <i>Human Molecular Genetics</i> , 2022, 31, 3133-3143.	1.4	11
485	<i>IGF2R</i> Missense Single-Nucleotide Polymorphisms and Breast Cancer Risk: The Multiethnic Cohort Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 1922-1924.	1.1	10
486	Genetic variation in the insulin, insulin-like growth factor, growth hormone, and leptin pathways in relation to breast cancer in African-American women: the AMBER consortium. <i>Npj Breast Cancer</i> , 2016, 2, .	2.3	10

#	ARTICLE	IF	CITATIONS
487	Genetic variants in the mTOR pathway and breast cancer risk in African American women. <i>Carcinogenesis</i> , 2016, 37, 49-55.	1.3	10
488	Atopic allergic conditions and pancreatic cancer risk: Results from the Multiethnic Cohort Study. <i>International Journal of Cancer</i> , 2018, 142, 2019-2027.	2.3	10
489	High mortality due to sepsis in Native Hawaiians and African Americans: The Multiethnic Cohort. <i>PLoS ONE</i> , 2017, 12, e0178374.	1.1	10
490	Meta-Analysis of Rare Variant Association Tests in Multiethnic Populations. <i>Genetic Epidemiology</i> , 2016, 40, 57-65.	0.6	9
491	Intake of cocoa products and risk of type-2 diabetes: the multiethnic cohort. <i>European Journal of Clinical Nutrition</i> , 2019, 73, 671-678.	1.3	9
492	Mendelian randomisation study of smoking exposure in relation to breast cancer risk. <i>British Journal of Cancer</i> , 2021, 125, 1135-1145.	2.9	9
493	A transcriptome-wide association study identifies novel candidate susceptibility genes for prostate cancer risk. <i>International Journal of Cancer</i> , 2022, 150, 80-90.	2.3	9
494	Trans-ethnic follow-up of breast cancer GWAS hits using the preferential linkage disequilibrium approach. <i>Oncotarget</i> , 2016, 7, 83160-83176.	0.8	9
495	<i>PHIP</i> - a novel candidate breast cancer susceptibility locus on 6q14.1. <i>Oncotarget</i> , 2017, 8, 102769-102782.	0.8	9
496	A Genome-Wide Pleiotropy Scan Does Not Identify New Susceptibility Loci for Estrogen Receptor Negative Breast Cancer. <i>PLoS ONE</i> , 2014, 9, e85955.	1.1	8
497	Genetic variation in the immunosuppression pathway genes and breast cancer susceptibility: a pooled analysis of 42,510 cases and 40,577 controls from the Breast Cancer Association Consortium. <i>Human Genetics</i> , 2016, 135, 137-154.	1.8	8
498	Validation of genetic associations with acute GVHD and nonrelapse mortality in DISCOVeRY-BMT. <i>Blood Advances</i> , 2019, 3, 2337-2341.	2.5	8
499	Variability in Cytogenetic Testing for Multiple Myeloma: A Comprehensive Analysis From Across the United States. <i>JCO Oncology Practice</i> , 2020, 16, e1169-e1180.	1.4	8
500	Genome-Wide Association Analyses Identify Variants in <i>IRF4</i> Associated With Acute Myeloid Leukemia and Myelodysplastic Syndrome Susceptibility. <i>Frontiers in Genetics</i> , 2021, 12, 554948.	1.1	8
501	Novel genetic variants associated with mortality after unrelated donor allogeneic hematopoietic cell transplantation. <i>EClinicalMedicine</i> , 2021, 40, 101093.	3.2	8
502	The Potential for Enhancing the Power of Genetic Association Studies in African Americans through the Reuse of Existing Genotype Data. <i>PLoS Genetics</i> , 2010, 6, e1001096.	1.5	8
503	Prognostic utility of self-reported sarcopenia (SARC) in the Multiethnic Cohort. <i>Journal of Cachexia, Sarcopenia and Muscle</i> , 2022, 13, 987-1002.	2.9	8
504	No evidence of interaction between known lipid-associated genetic variants and smoking in the multi-ethnic PAGE population. <i>Human Genetics</i> , 2013, 132, 1427-1431.	1.8	7

#	ARTICLE	IF	CITATIONS
505	Fine-Mapping <i>IGF1</i> and Prostate Cancer Risk in African Americans: The Multiethnic Cohort Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1928-1932.	1.1	7
506	Diet quality measured by four a priori-defined diet quality indices is associated with lipid-soluble micronutrients in the Multiethnic Cohort Study (MEC). <i>European Journal of Clinical Nutrition</i> , 2019, 73, 703-713.	1.3	7
507	Multiple functional variants in the <i>IL1RL1</i> region are pretransplant markers for risk of GVHD and infection deaths. <i>Blood Advances</i> , 2019, 3, 2512-2524.	2.5	7
508	Association of germline genetic variants with breast cancer-specific survival in patient subgroups defined by clinic-pathological variables related to tumor biology and type of systemic treatment. <i>Breast Cancer Research</i> , 2021, 23, 86.	2.2	7
509	Combined Donor and Recipient Non-HLA Genotypes Show Evidence of Genome Wide Association with Transplant Related Mortality (TRM) after HLA-Matched Unrelated Donor Blood and Marrow Transplantation (URD-BMT) (DISCOVeRY-BMT study). <i>Blood</i> , 2015, 126, 61-61.	0.6	7
510	Modifying effects of race and ethnicity and <i>APOE</i> on the association of physical activity with risk of Alzheimer's disease and related dementias. <i>Alzheimer's and Dementia</i> , 2023, 19, 507-517.	0.4	7
511	Genetic variation in the <i>HSD17B1</i> gene and risk of prostate cancer. <i>PLoS Genetics</i> , 2005, preprint, e68.	1.5	6
512	One thousand genomes imputation in the national cancer institute breast and prostate cancer cohort consortium aggressive prostate cancer genome-wide association study. <i>Prostate</i> , 2013, 73, 677-689.	1.2	6
513	Deep targeted sequencing of 12 breast cancer susceptibility regions in 4611 women across four different ethnicities. <i>Breast Cancer Research</i> , 2016, 18, 109.	2.2	6
514	Complex patterns of direct and indirect association between the transcription Factor-7 like 2 gene, body mass index and type 2 diabetes diagnosis in adulthood in the Hispanic Community Health Study/Study of Latinos. <i>BMC Obesity</i> , 2018, 5, 26.	3.1	6
515	Association between mitochondrial genetic variation and breast cancer risk: The Multiethnic Cohort. <i>PLoS ONE</i> , 2019, 14, e0222284.	1.1	6
516	Evaluation of vitamin D biosynthesis and pathway target genes reveals <i>UGT2A1/2</i> and <i>EGFR</i> polymorphisms associated with epithelial ovarian cancer in African American Women. <i>Cancer Medicine</i> , 2019, 8, 2503-2513.	1.3	6
517	Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. <i>American Journal of Human Genetics</i> , 2021, 108, 1190-1203.	2.6	6
518	Large-scale cross-cancer fine-mapping of the 5p15.33 region reveals multiple independent signals. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100041.	1.0	6
519	Novel strategy for disease risk prediction incorporating predicted gene expression and DNA methylation data: a multi-phased study of prostate cancer. <i>Cancer Communications</i> , 2021, 41, 1387-1397.	3.7	6
520	Pre-HCT mosaicism increases relapse risk and lowers survival in acute lymphoblastic leukemia patients post-unrelated HCT. <i>Blood Advances</i> , 2021, 5, 66-70.	2.5	6
521	Rare germline copy number variants (CNVs) and breast cancer risk. <i>Communications Biology</i> , 2022, 5, 65.	2.0	6
522	Enrichment analyses identify shared associations for 25 quantitative traits in over 600,000 individuals from seven diverse ancestries. <i>American Journal of Human Genetics</i> , 2022, 109, 871-884.	2.6	6

#	ARTICLE	IF	CITATIONS
523	A Genome-Wide Gene-Based Gene-Environment Interaction Study of Breast Cancer in More than 90,000 Women. <i>Cancer Research Communications</i> , 2022, 2, 211-219.	0.7	6
524	Post-Genome-Wide Association Study Challenges for Lipid Traits: Describing Age as a Modifier of Gene-Lipid Associations in the Population Architecture Using Genomics and Epidemiology (PAGE) Study. <i>Annals of Human Genetics</i> , 2013, 77, 416-425.	0.3	5
525	Stepwise approach to SNP-set analysis illustrated with the MetaboChip and colorectal cancer in Japanese Americans of the Multiethnic Cohort. <i>BMC Genomics</i> , 2018, 19, 524.	1.2	5
526	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , 2019, 9, 12524.	1.6	5
527	CYP3A7*1C allele: linking premenopausal oestrogen and progesterone levels with risk of hormone receptor-positive breast cancers. <i>British Journal of Cancer</i> , 2021, 124, 842-854.	2.9	5
528	KLK3 SNP-SNP interactions for prediction of prostate cancer aggressiveness. <i>Scientific Reports</i> , 2021, 11, 9264.	1.6	5
529	Evidence for Heterogeneous Genetic Associations with Acute Lymphoblastic Leukemia (ALL) By Cytogenetics and Sex in High-Risk Patients Treated with Matched Unrelated Donor Allogeneic Blood or Marrow Transplant (URD-BMT). <i>Blood</i> , 2015, 126, 2621-2621.	0.6	5
530	Contributions of Social Factors to Disparities in Prostate Cancer Risk Profiles among Black Men and Non-Hispanic White Men with Prostate Cancer in California. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 404-412.	1.1	5
531	Predicted gene expression in ancestrally diverse populations leads to discovery of susceptibility loci for lifestyle and cardiometabolic traits. <i>American Journal of Human Genetics</i> , 2022, 109, 669-679.	2.6	5
532	Germ Line Variation at 8q24 and Endometrial Cancer Risk: Table 1.. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 2166-2168.	1.1	4
533	Tobacco smoke toxicant and carcinogen biomarkers and lung cancer susceptibility in smokers. <i>Journal of Thoracic Oncology</i> , 2016, 11, S7-S8.	0.5	4
534	ED01.02 Tobacco Carcinogens and Lung Cancer Susceptibility. <i>Journal of Thoracic Oncology</i> , 2017, 12, S19-S20.	0.5	4
535	Association of Combined Sero-Positivity to <i>Helicobacter pylori</i> and <i>Streptococcus gallolyticus</i> with Risk of Colorectal Cancer. <i>Microorganisms</i> , 2020, 8, 1698.	1.6	4
536	Gene-Environment Interactions Relevant to Estrogen and Risk of Breast Cancer: Can Gene-Environment Interactions Be Detected Only among Candidate SNPs from Genome-Wide Association Studies?. <i>Cancers</i> , 2021, 13, 2370.	1.7	4
537	Prognostic impact of pre-transplant chromosomal aberrations in peripheral blood of patients undergoing unrelated donor hematopoietic cell transplant for acute myeloid leukemia. <i>Scientific Reports</i> , 2021, 11, 15004.	1.6	4
538	Discovery of structural deletions in breast cancer predisposition genes using whole genome sequencing data from 2000 women of African-ancestry. <i>Human Genetics</i> , 2021, 140, 1449-1457.	1.8	4
539	The efficacy of detecting variants with small effects on the Affymetrix 6.0 platform using pooled DNA. <i>Human Genetics</i> , 2011, 130, 607-621.	1.8	3
540	The Impact of GWAS Findings on Cancer Etiology and Prevention. <i>Current Epidemiology Reports</i> , 2014, 1, 130-137.	1.1	3

#	ARTICLE	IF	CITATIONS
541	AA9int: SNP interaction pattern search using non-hierarchical additive model set. <i>Bioinformatics</i> , 2018, 34, 4141-4150.	1.8	3
542	Alcohol Intake and Alcoholâ€“SNP Interactions Associated with Prostate Cancer Aggressiveness. <i>Journal of Clinical Medicine</i> , 2021, 10, 553.	1.0	3
543	Prediagnostic Antibody Responses to <i>Fusobacterium nucleatum</i> Proteins Are Not Associated with Risk of Colorectal Cancer in a Large U.S. Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 1279-1282.	1.1	3
544	Ancestral diversity improves discovery and fine-mapping of genetic loci for anthropometric traitsâ€“The Hispanic/Latino Anthropometry Consortium. <i>Human Genetics and Genomics Advances</i> , 2022, 3, 100099.	1.0	3
545	Cancer Mortality Patterns by Birthplace and Generation Status of Mexican Latinos: The Multiethnic Cohort. <i>Journal of the National Cancer Institute</i> , 2022, 114, 959-968.	3.0	3
546	Methodological Considerations in Estimation of Phenotype Heritability Using Genome-Wide SNP Data, Illustrated by an Analysis of the Heritability of Height in a Large Sample of African Ancestry Adults. <i>PLoS ONE</i> , 2015, 10, e0131106.	1.1	2
547	rs2735383, located at a microRNA binding site in the 3â€™UTR of NBS1, is not associated with breast cancer risk. <i>Scientific Reports</i> , 2016, 6, 36874.	1.6	2
548	Interactions between breast cancer susceptibility loci and menopausal hormone therapy in relationship to breast cancer in the Breast and Prostate Cancer Cohort Consortium. <i>Breast Cancer Research and Treatment</i> , 2016, 155, 531-540.	1.1	2
549	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , 2020, 10, 9688.	1.6	2
550	Association of Anthropometric Measures With the Risk of Prostate Cancer in the Multiethnic Cohort. <i>American Journal of Epidemiology</i> , 2021, 190, 1770-1783.	1.6	2
551	Genome-wide association study of pancreatic fat: The Multiethnic Cohort Adiposity Phenotype Study. <i>PLoS ONE</i> , 2021, 16, e0249615.	1.1	2
552	Cholesterol lowering drug use and breast cancer survival: the Multiethnic Cohort Study. <i>Breast Cancer Research and Treatment</i> , 2021, 190, 165-173.	1.1	2
553	Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. <i>Scientific Reports</i> , 2021, 11, 19787.	1.6	2
554	A Meta-Analysis Of Genome-Wide Association Studies Of Multiple Myeloma In Cases and Controls Of European Origin Identifies a Risk Locus In 12q23.1. <i>Blood</i> , 2013, 122, 3111-3111.	0.6	2
555	De Novo and Therapy-Related Acute Myeloid Leukemia and Myelodysplastic Syndrome: Similarities and Differences in SNP-Array Detected Chromosomal Aberrations in Pre-Transplant Blood Samples. <i>Blood</i> , 2019, 134, 1430-1430.	0.6	2
556	Change in the inflammatory potential of diet over 10 years and subsequent mortality: the Multiethnic Cohort Study. <i>British Journal of Nutrition</i> , 2022, , 1-23.	1.2	2
557	Evaluating breast cancer predisposition genes in women of African ancestry. <i>Genetics in Medicine</i> , 2022, 24, 1468-1475.	1.1	2
558	Genome-wide interaction analysis of menopausal hormone therapy use and breast cancer risk among 62,370 women. <i>Scientific Reports</i> , 2022, 12, 6199.	1.6	2

#	ARTICLE	IF	CITATIONS
559	Genetic Risk of Second Primary Cancer in Breast Cancer Survivors: The Multiethnic Cohort Study. <i>Cancer Research</i> , 2022, 82, 3201-3208.	0.4	2
560	Cross-Cancer Analysis Reveals Novel Pleiotropic Associationsâ€™Response. <i>Cancer Research</i> , 2017, 77, 6045-6046.	0.4	1
561	Data-adaptive multi-locus association testing in subjects with arbitrary genealogical relationships. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2019, 18, .	0.2	1
562	Genetic variants in anti-MÃ¼llerian hormone-related genes and breast cancer risk: results from the AMBER consortium. <i>Breast Cancer Research and Treatment</i> , 2021, 185, 469-478.	1.1	1
563	Explaining the Excess Risk of Multiple Myeloma in African-Americans. <i>Blood</i> , 2012, 120, 4002-4002.	0.6	1
564	HLA Haplotypes Are Associated with Multiple Myeloma Risk in the African American Multiple Myeloma Study (AAMMS). <i>Blood</i> , 2016, 128, 3250-3250.	0.6	1
565	Clonal Hematopoiesis with Somatic Mutations Is a Common, Age-Related Condition Associated with Adverse Outcomes. <i>Blood</i> , 2014, 124, 840-840.	0.6	1
566	Genome-Wide Association Study of Overall and Progression-Free Survival after HLA-Matched Unrelated Donor Blood and Marrow Transplantation (DISCOVeRY-BMT study). <i>Blood</i> , 2015, 126, 397-397.	0.6	1
567	Urinary 6-sulfatoxymelatonin Levels and Prostate Cancer Risk among Men in the Multiethnic Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 688-691.	1.1	1
568	Improved Imputation of Common and Uncommon Single Nucleotide Polymorphisms (SNPs) with a New Reference Set. <i>Nature Precedings</i> , 2011, , .	0.1	0
569	A comprehensive analysis of polymorphic variants in steroid hormone and insulinâ€™like growth factorâ€™1 metabolism and risk of <i>in situ</i> breast cancer: Results from the Breast and Prostate Cancer Cohort Consortium. <i>International Journal of Cancer</i> , 2018, 142, 1182-1188.	2.3	0
570	Genetic Associations with Day +100 Transplant Related Mortality (TRM) after HLA-Matched Unrelated Donor (MUD) Blood and Marrow Transplantation (DISCOVeRY-BMT Study). <i>Biology of Blood and Marrow Transplantation</i> , 2018, 24, S81-S82.	2.0	0
571	White Rice Consumption and Risk for Colorectal Cancer among Japanese Americans: The Multiethnic Cohort Study. <i>Journal of Epidemiology</i> , 2021, , .	1.1	0
572	Abstract LB011: Meta-analysis in more than 80,000 men of African ancestry identified nine novel variants associated with prostate cancer. , 2021, , .		0
573	A High-Density Admixture Scan in 1,670 African Americans with Hypertension. <i>PLoS Genetics</i> , 2005, preprint, e196.	1.5	0
574	Polymorphisms In IRS1 and IL6R and Susceptibility To Multiple Myeloma. <i>Blood</i> , 2013, 122, 3154-3154.	0.6	0
575	Obesity In Young Adulthood Is Associated With Early Onset Multiple Myeloma In African Americans. <i>Blood</i> , 2013, 122, 1872-1872.	0.6	0
576	Genetic Susceptibility Markers of Multiple Myeloma in African-Americans. <i>Blood</i> , 2014, 124, 2030-2030.	0.6	0

#	ARTICLE	IF	CITATIONS
577	Replication of Candidate SNP Survival Analyses and Gene-Based Tests of Association with Survival Outcomes after an Unrelated Donor Blood or Marrow Transplant: Results from the Discovery-BMT Study. <i>Blood</i> , 2016, 128, 71-71.	0.6	0
578	Exome Array Analyses Identify New Genes Influencing Survival Outcomes after HLA-Matched Unrelated Donor Blood and Marrow Transplantation. <i>Blood</i> , 2016, 128, 518-518.	0.6	0
579	Exome Array Analyses Identify Low-Frequency Germline Variants Associated with Increased Risk of AML in a HLA-Matched Unrelated Donor Blood and Marrow Transplant Population. <i>Blood</i> , 2016, 128, 42-42.	0.6	0
580	Genome Wide Association Analyses Identify Pleiotropic Variants Associated with Acute Myeloid Leukemia (AML) and Myelodysplastic Syndrome (MDS) Susceptibility. <i>Blood</i> , 2018, 132, 1500-1500.	0.6	0
581	Multiple Functional Donor Polymorphisms in IL1RL1 region Associate with Death Due to GvHD or Infection after Unrelated Donor Allogeneic Hematopoietic Stem Cell Transplantation (HCT) for AML and MDS. <i>Blood</i> , 2018, 132, 312-312.	0.6	0
582	Comprehensive Investigation of White Blood Cell and Gene Expression Profiles As Risk Factors for Multiple Myeloma in African Americans. <i>Blood</i> , 2019, 134, 4379-4379.	0.6	0
583	Genome Wide Interaction Analysis Identifies Expression Quantitative Trait Loci Associated with Reduced Survival after Reduced Intensity Conditioning HLA-Matched Unrelated Donor Allogeneic Hematopoietic Cell Transplant. <i>Blood</i> , 2019, 134, 4595-4595.	0.6	0
584	Meta-Analysis of Genome-Wide Association Studies of Acute Myeloid Leukemia (AML) Patients Identifies Variants Associated with Risk of 11q23/KMT2A-Translocated and Core-Binding Factor (CBF) AML and Suggests a Role for Transcription Elongation in Leukemogenesis. <i>Blood</i> , 2020, 136, 29-30.	0.6	0
585	Population Distribution of GvL and GvH Minor Histocompatibility Antigens. <i>Blood</i> , 2020, 136, 23-25.	0.6	0
586	Associations of Clinical Outcomes after Allogeneic Hematopoietic Cell Transplantation with Number of Predicted Class II Restricted mHA. <i>Blood</i> , 2020, 136, 2-2.	0.6	0
587	Characteristics of and Risk Factors for Monoclonal Gammopathy of Undetermined Significance (MGUS) in the Multiethnic Cohort Study. <i>Blood</i> , 2020, 136, 28-29.	0.6	0
588	Pre-Transplant Clonal Mosaicism Is Associated with Increased Relapse and Lower Survival in Acute Lymphoblastic Leukemia Patients Undergoing Allogeneic Hematopoietic Cell Transplant. <i>Blood</i> , 2020, 136, 9-10.	0.6	0
589	Abstract P2-09-01: Population-based risk estimates of clinical subtypes of breast cancer among carriers of germline pathogenic variants in cancer predisposition genes. <i>Cancer Research</i> , 2022, 82, P2-09-01-P2-09-01.	0.4	0
590	The Association of Prediagnostic Statin Use with Aggressive Prostate Cancer from the Multiethnic Cohort Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2022, 31, 999-1005.	1.1	0