

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

140
papers

10,975
citations

54
h-index

104
g-index

150
ext. papers

14,215
ext. citations

9.1
avg, IF

5.28
L-index

#	Paper	IF	Citations
140	The Immune Landscape of Cancer. <i>Immunity</i> , 2018 , 48, 812-830.e14	32.3	1754
139	The importance of race and ethnic background in biomedical research and clinical practice. <i>New England Journal of Medicine</i> , 2003 , 348, 1170-5	59.2	813
138	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017 , 551, 92-94	50.4	643
137	Association of common genetic variation in the insulin/IGF1 signaling pathway with human longevity. <i>Aging Cell</i> , 2009 , 8, 460-72	9.9	279
136	Reduced neutrophil count in people of African descent is due to a regulatory variant in the Duffy antigen receptor for chemokines gene. <i>PLoS Genetics</i> , 2009 , 5, e1000360	6	251
135	Latino populations: a unique opportunity for the study of race, genetics, and social environment in epidemiological research. <i>American Journal of Public Health</i> , 2005 , 95, 2161-8	5.1	251
134	Population stratification confounds genetic association studies among Latinos. <i>Human Genetics</i> , 2006 , 118, 652-64	6.3	207
133	Prevention of breast cancer in postmenopausal women: approaches to estimating and reducing risk. <i>Journal of the National Cancer Institute</i> , 2009 , 101, 384-98	9.7	199
132	Pharmacogenetic differences in response to albuterol between Puerto Ricans and Mexicans with asthma. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2005 , 171, 563-70	10.2	198
131	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017 , 49, 1767-1778	36.3	186
130	Genetic ancestry in lung-function predictions. <i>New England Journal of Medicine</i> , 2010 , 363, 321-30	59.2	182
129	Mammographic breast density and the Gail model for breast cancer risk prediction in a screening population. <i>Breast Cancer Research and Treatment</i> , 2005 , 94, 115-22	4.4	182
128	Development of a panel of genome-wide ancestry informative markers to study admixture throughout the Americas. <i>PLoS Genetics</i> , 2012 , 8, e1002554	6	176
127	Genome-wide meta-analysis points to CTC1 and ZNF676 as genes regulating telomere homeostasis in humans. <i>Human Molecular Genetics</i> , 2012 , 21, 5385-94	5.6	162
126	Lower bronchodilator responsiveness in Puerto Rican than in Mexican subjects with asthma. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2004 , 169, 386-92	10.2	153
125	Admixture mapping of white cell count: genetic locus responsible for lower white blood cell count in the Health ABC and Jackson Heart studies. <i>American Journal of Human Genetics</i> , 2008 , 82, 81-7	11	150
124	Genetic determinants of serum testosterone concentrations in men. <i>PLoS Genetics</i> , 2011 , 7, e1002313	6	148

123	Admixture mapping of an allele affecting interleukin 6 soluble receptor and interleukin 6 levels. <i>American Journal of Human Genetics</i> , 2007 , 80, 716-26	11	145
122	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, 5373-84	5.6	143
121	The contributions of breast density and common genetic variation to breast cancer risk. <i>Journal of the National Cancer Institute</i> , 2015 , 107,	9.7	128
120	CD36 repression activates a multicellular stromal program shared by high mammographic density and tumor tissues. <i>Cancer Discovery</i> , 2012 , 2, 826-39	24.4	128
119	Population structure, admixture, and aging-related phenotypes in African American adults: the Cardiovascular Health Study. <i>American Journal of Human Genetics</i> , 2005 , 76, 463-77	11	127
118	ADAM33 is not associated with asthma in Puerto Rican or Mexican populations. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2003 , 168, 1312-6	10.2	120
117	Genome-wide association study of white blood cell count in 16,388 African Americans: the continental origins and genetic epidemiology network (COGENT). <i>PLoS Genetics</i> , 2011 , 7, e1002108	6	111
116	Cancer health disparities in racial/ethnic minorities in the United States. <i>British Journal of Cancer</i> , 2021 , 124, 315-332	8.7	110
115	Mammographic breast density and family history of breast cancer. <i>Journal of the National Cancer Institute</i> , 2003 , 95, 556-8	9.7	104
114	Leveraging genetic variability across populations for the identification of causal variants. <i>American Journal of Human Genetics</i> , 2010 , 86, 23-33	11	103
113	Breast Cancer Screening in the Precision Medicine Era: Risk-Based Screening in a Population-Based Trial. <i>Journal of the National Cancer Institute</i> , 2017 , 109,	9.7	102
112	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	36.3	101
111	Heterogeneity in genetic admixture across different regions of Argentina. <i>PLoS ONE</i> , 2012 , 7, e34695	3.7	100
110	Race and Genetic Ancestry in Medicine - A Time for Reckoning with Racism. <i>New England Journal of Medicine</i> , 2021 , 384, 474-480	59.2	100
109	Human population structure and genetic association studies. <i>Pharmacogenomics</i> , 2003 , 4, 431-41	2.6	97
108	Association between the T29-->C polymorphism in the transforming growth factor beta1 gene and breast cancer among elderly white women: The Study of Osteoporotic Fractures. <i>JAMA - Journal of the American Medical Association</i> , 2001 , 285, 2859-63	27.4	93
107	CD14 tobacco gene-environment interaction modifies asthma severity and immunoglobulin E levels in Latinos with asthma. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2005 , 172, 173-82	10.2	91
106	Genome-wide association study of breast cancer in Latinas identifies novel protective variants on 6q25. <i>Nature Communications</i> , 2014 , 5, 5260	17.4	89

105	Genetic ancestry and risk of breast cancer among U.S. Latinas. <i>Cancer Research</i> , 2008 , 68, 9723-8	10.1	89
104	Comparison of three methods to estimate genetic ancestry and control for stratification in genetic association studies among admixed populations. <i>Human Genetics</i> , 2005 , 118, 424-33	6.3	89
103	The genetics of human longevity. <i>American Journal of Medicine</i> , 2004 , 117, 851-60	2.4	88
102	A meta-analysis and genome-wide association study of platelet count and mean platelet volume in african americans. <i>PLoS Genetics</i> , 2012 , 8, e1002491	6	84
101	Breast cancer risk prediction using a clinical risk model and polygenic risk score. <i>Breast Cancer Research and Treatment</i> , 2016 , 159, 513-25	4.4	82
100	A review of cancer in U.S. Hispanic populations. <i>Cancer Prevention Research</i> , 2012 , 5, 150-63	3.2	81
99	European ancestry is positively associated with breast cancer risk in Mexican women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010 , 19, 1074-82	4	80
98	Clinical and biomarker predictors of side effects from tamoxifen. <i>Breast Cancer Research and Treatment</i> , 2012 , 132, 1107-18	4.4	78
97	Are breast density and bone mineral density independent risk factors for breast cancer?. <i>Journal of the National Cancer Institute</i> , 2005 , 97, 368-74	9.7	73
96	History shaped the geographic distribution of genomic admixture on the island of Puerto Rico. <i>PLoS ONE</i> , 2011 , 6, e16513	3.7	71
95	Early therapy for latent tuberculosis infection. <i>American Journal of Epidemiology</i> , 2001 , 153, 381-5	3.8	69
94	Genetic ancestry, population sub-structure, and cardiovascular disease-related traits among African-American participants in the CARDIA Study. <i>Human Genetics</i> , 2007 , 121, 565-75	6.3	68
93	Whole-Genome Sequencing of Pharmacogenetic Drug Response in Racially Diverse Children with Asthma. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2018 , 197, 1552-1564	10.2	65
92	Genome-wide screen for asthma in Puerto Ricans: evidence for association with 5q23 region. <i>Human Genetics</i> , 2008 , 123, 455-68	6.3	63
91	Genetic variation in insulin/IGF-1 signaling pathways and longevity. <i>Ageing Research Reviews</i> , 2011 , 10, 201-4	12	61
90	Mammographic density and estrogen receptor status of breast cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2004 , 13, 2090-5	4	57
89	Comprehensive Analysis of Genetic Ancestry and Its Molecular Correlates in Cancer. <i>Cancer Cell</i> , 2020 , 37, 639-654.e6	24.3	56
88	Serum insulin-like growth factor-1 binding proteins 1 and 2 and mortality in older adults: the Health, Aging, and Body Composition Study. <i>Journal of the American Geriatrics Society</i> , 2009 , 57, 1213-8	5.6	55

87	Ancestry-environment interactions and asthma risk among Puerto Ricans. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2006 , 174, 1088-93	10.2	54
86	Tracing sub-structure in the European American population with PCA-informative markers. <i>PLoS Genetics</i> , 2008 , 4, e1000114	6	53
85	Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2019 , 48, 795-806	7.8	52
84	Genetic ancestry and risk factors for breast cancer among Latinas in the San Francisco Bay Area. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006 , 15, 1878-85	4	51
83	Admixture mapping identifies a locus on 6q25 associated with breast cancer risk in US Latinas. <i>Human Molecular Genetics</i> , 2012 , 21, 1907-17	5.6	50
82	Genome-wide association analysis of red blood cell traits in African Americans: the COGENT Network. <i>Human Molecular Genetics</i> , 2013 , 22, 2529-38	5.6	48
81	Potential public health impact of new tuberculosis vaccines. <i>Emerging Infectious Diseases</i> , 2004 , 10, 1529-35	9.5	47
80	Epigenetic and transcriptional determinants of the human breast. <i>Nature Communications</i> , 2015 , 6, 6351	17.4	44
79	The association between a Darc gene polymorphism and clinical outcomes in African American patients with acute lung injury. <i>Chest</i> , 2012 , 141, 1160-1169	5.3	44
78	Estrogen receptor alpha haplotypes and breast cancer risk in older Caucasian women. <i>Breast Cancer Research and Treatment</i> , 2007 , 106, 273-80	4.4	44
77	Inflammation and stress-related candidate genes, plasma interleukin-6 levels, and longevity in older adults. <i>Experimental Gerontology</i> , 2009 , 44, 350-5	4.5	43
76	An admixture scan in 1,484 African American women with breast cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009 , 18, 3110-7	4	42
75	Duffy (Fy), DARC, and neutropenia among women from the United States, Europe and the Caribbean. <i>British Journal of Haematology</i> , 2008 , 143, 288-93	4.5	42
74	Risk Factors That Increase Risk of Estrogen Receptor-Positive and -Negative Breast Cancer. <i>Journal of the National Cancer Institute</i> , 2017 , 109,	9.7	39
73	Admixture-matched case-control study: a practical approach for genetic association studies in admixed populations. <i>Human Genetics</i> , 2006 , 118, 626-39	6.3	39
72	African ancestry, socioeconomic status, and kidney function in elderly African Americans: a genetic admixture analysis. <i>Journal of the American Society of Nephrology: JASN</i> , 2006 , 17, 3491-6	12.7	37
71	Pan-cancer study detects genetic risk variants and shared genetic basis in two large cohorts. <i>Nature Communications</i> , 2020 , 11, 4423	17.4	36
70	Methylenetetrahydrofolate reductase C677T polymorphism and cognitive function in older women. <i>American Journal of Epidemiology</i> , 2007 , 166, 672-8	3.8	32

69	Genome-wide association study identifies variants at 16p13 associated with survival in multiple myeloma patients. <i>Nature Communications</i> , 2015 , 6, 7539	17.4	31
68	Genetic admixture, adipocytokines, and adiposity in Black Americans: the Health, Aging, and Body Composition study. <i>Human Genetics</i> , 2007 , 121, 615-24	6.3	31
67	Genomic Profiling of Prostate Cancers from Men with African and European Ancestry. <i>Clinical Cancer Research</i> , 2020 , 26, 4651-4660	12.9	31
66	The Association of African Ancestry and elevated creatinine in the Coronary Artery Risk Development in Young Adults (CARDIA) Study. <i>American Journal of Nephrology</i> , 2010 , 31, 202-8	4.6	30
65	Cis-eQTL-based trans-ethnic meta-analysis reveals novel genes associated with breast cancer risk. <i>PLoS Genetics</i> , 2017 , 13, e1006690	6	29
64	Genetic ancestry and risk of mortality among U.S. Latinas with breast cancer. <i>Cancer Research</i> , 2013 , 73, 7243-53	10.1	28
63	Germline genetic contribution to the immune landscape of cancer. <i>Immunity</i> , 2021 , 54, 367-386.e8	32.3	27
62	Association of estrogen receptor alpha polymorphisms with breast cancer risk in older Caucasian women. <i>International Journal of Cancer</i> , 2005 , 116, 984-91	7.5	25
61	Genetic ancestry-smoking interactions and lung function in African Americans: a cohort study. <i>PLoS ONE</i> , 2012 , 7, e39541	3.7	25
60	Recurrent BRCA1 and BRCA2 mutations in Mexican women with breast cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 498-505	4	24
59	The genetics of benign neutropenia. <i>Israel Medical Association Journal</i> , 2011 , 13, 625-9	0.9	24
58	Identification of novel common breast cancer risk variants at the 6q25 locus among Latinas. <i>Breast Cancer Research</i> , 2019 , 21, 3	8.3	23
57	Genetic ancestry modifies the association between genetic risk variants and breast cancer risk among Hispanic and non-Hispanic white women. <i>Carcinogenesis</i> , 2013 , 34, 1787-93	4.6	22
56	A genome-wide association study identifies only two ancestry specific variants associated with spontaneous preterm birth. <i>Scientific Reports</i> , 2018 , 8, 226	4.9	21
55	Common variants in the CRP gene in relation to longevity and cause-specific mortality in older adults: the Cardiovascular Health Study. <i>Atherosclerosis</i> , 2008 , 197, 922-30	3.1	21
54	Estrogen metabolizing polymorphisms and breast cancer risk among older white women. <i>Breast Cancer Research and Treatment</i> , 2005 , 93, 261-70	4.4	21
53	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	9.7	21
52	Genetic Investigation Into the Differential Risk of Atrial Fibrillation Among Black and White Individuals. <i>JAMA Cardiology</i> , 2016 , 1, 442-50	16.2	20

51	Pathogenic and likely pathogenic variants in PALB2, CHEK2, and other known breast cancer susceptibility genes among 1054 BRCA-negative Hispanics with breast cancer. <i>Cancer</i> , 2019 , 125, 2829-2836	6.4	19
50	Association between rs2981582 polymorphism in the FGFR2 gene and the risk of breast cancer in Mexican women. <i>Archives of Medical Research</i> , 2013 , 44, 459-66	6.6	19
49	Pan-cancer analysis demonstrates that integrating polygenic risk scores with modifiable risk factors improves risk prediction. <i>Nature Communications</i> , 2020 , 11, 6084	17.4	19
48	Thyroid hormones and breast cancer association according to menopausal status and body mass index. <i>Breast Cancer Research</i> , 2018 , 20, 94	8.3	18
47	No association between the TGF-beta 1 Leu10Pro polymorphism and osteoporosis among white women in the United States. <i>American Journal of Medicine</i> , 2003 , 114, 227-31	2.4	18
46	Family History and Breast Cancer Risk Among Older Women in the Breast Cancer Surveillance Consortium Cohort. <i>JAMA Internal Medicine</i> , 2018 , 178, 494-501	11.5	17
45	Telomere length and the risk of atrial fibrillation: insights into the role of biological versus chronological aging. <i>Circulation: Arrhythmia and Electrophysiology</i> , 2014 , 7, 1026-32	6.4	16
44	Identification of Pleiotropic Cancer Susceptibility Variants from Genome-Wide Association Studies Reveals Functional Characteristics. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018 , 27, 75-85	4	16
43	Joint relative risks for estrogen receptor-positive breast cancer from a clinical model, polygenic risk score, and sex hormones. <i>Breast Cancer Research and Treatment</i> , 2017 , 166, 603-612	4.4	14
42	Using Breast Cancer Risk Associated Polymorphisms to Identify Women for Breast Cancer Chemoprevention. <i>PLoS ONE</i> , 2017 , 12, e0168601	3.7	14
41	Comparing Ethnicity-Specific Reference Intervals for Clinical Laboratory Tests from EHR Data. <i>Journal of Applied Laboratory Medicine</i> , 2018 , 3, 366-377	2	14
40	Multiple breast cancer risk variants are associated with differential transcript isoform expression in tumors. <i>Human Molecular Genetics</i> , 2015 , 24, 7421-31	5.6	13
39	Association of imputed prostate cancer transcriptome with disease risk reveals novel mechanisms. <i>Nature Communications</i> , 2019 , 10, 3107	17.4	13
38	Estimation of tamoxifen metabolite concentrations in the blood of breast cancer patients through CYP2D6 genotype activity score. <i>Breast Cancer Research and Treatment</i> , 2012 , 133, 677-83	4.4	13
37	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	4	13
36	A response to "Personalised medicine and population health: breast and ovarian cancer". <i>Human Genetics</i> , 2019 , 138, 287-289	6.3	13
35	Interaction between common breast cancer susceptibility variants, genetic ancestry, and nongenetic risk factors in Hispanic women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 1731-8	4	12
34	Intrahepatic Cholestasis of Pregnancy (ICP) in U.S. Latinas and Chileans: Clinical features, Ancestry Analysis, and Admixture Mapping. <i>PLoS ONE</i> , 2015 , 10, e0131211	3.7	11

33	Moderate-intensity physical activity ameliorates the breast cancer risk in diabetic women. <i>Diabetes Care</i> , 2012 , 35, 2500-2	14.6	10
32	Human Epidermal Growth Factor Receptor 2-Positive Breast Cancer Is Associated with Indigenous American Ancestry in Latin American Women. <i>Cancer Research</i> , 2020 , 80, 1893-1901	10.1	10
31	Genetic Variation in CCL5 Signaling Genes and Triple Negative Breast Cancer: Susceptibility and Prognosis Implications. <i>Frontiers in Oncology</i> , 2019 , 9, 1328	5.3	9
30	Coinherited genetics of multiple myeloma and its precursor, monoclonal gammopathy of undetermined significance. <i>Blood Advances</i> , 2020 , 4, 2789-2797	7.8	8
29	Immunotherapy-Mediated Thyroid Dysfunction: Genetic Risk and Impact on Outcomes with PD-1 Blockade in Non-Small Cell Lung Cancer. <i>Clinical Cancer Research</i> , 2021 , 27, 5131-5140	12.9	7
28	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	7.8	5
27	European genetic ancestry associated with risk of childhood ependymoma. <i>Neuro-Oncology</i> , 2020 , 22, 1637-1646	1	5
26	Assessing the performance of in silico methods for predicting the pathogenicity of variants in the gene CHEK2, among Hispanic females with breast cancer. <i>Human Mutation</i> , 2019 , 40, 1612-1622	4.7	4
25	Ancestry-associated transcriptomic profiles of breast cancer in patients of African, Arab, and European ancestry. <i>Npj Breast Cancer</i> , 2021 , 7, 10	7.8	4
24	Lung Function in African American Children with Asthma Is Associated with Novel Regulatory Variants of the KIT Ligand and Gene-By-Air-Pollution Interaction. <i>Genetics</i> , 2020 , 215, 869-886	4	3
23	The CYP17A1 -34T > C polymorphism and breast cancer risk in BRCA1 and BRCA2 mutation carriers. <i>Breast Cancer Research and Treatment</i> , 2011 , 126, 521-7	4.4	3
22	Recent shifts in the genomic ancestry of Mexican Americans may alter the genetic architecture of biomedical traits. <i>ELife</i> , 2020 , 9,	8.9	3
21	Two truncating variants in FANCC and breast cancer risk. <i>Scientific Reports</i> , 2019 , 9, 12524	4.9	2
20	Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. <i>Scientific Reports</i> , 2020 , 10, 9688	4.9	2
19	Evaluation of the association of heterozygous germline variants in NTHL1 with breast cancer predisposition: an international multi-center study of 47,180 subjects. <i>Npj Breast Cancer</i> , 2021 , 7, 52	7.8	2
18	Whole-Genome Sequencing Identifies Novel Functional Loci Associated with Lung Function in Puerto Rican Youth. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020 , 202, 962-972	10.2	1
17	Interval breast cancers - insights into a complex phenotype. <i>Nature Reviews Clinical Oncology</i> , 2020 , 17, 138-139	19.4	1
16	Energy homeostasis genes modify the association between serum concentrations of IGF-1 and IGFBP-3 and breast cancer risk.. <i>Scientific Reports</i> , 2022 , 12, 1837	4.9	1

15	Systems Biology for Multiplatform Data Integration: An Overview. <i>Methods in Molecular Biology</i> , 2020 , 2055, 641-647	1.4	1
14	A polygenic risk score for breast cancer in U.S. Latinas and Latin-American women		1
13	Expression quantitative trait loci of genes predicting outcome are associated with survival of multiple myeloma patients. <i>International Journal of Cancer</i> , 2021 , 149, 327-336	7.5	1
12	Cross-ancestry GWAS meta-analysis identifies six breast cancer loci in African and European ancestry women. <i>Nature Communications</i> , 2021 , 12, 4198	17.4	1
11	Genetics of breast cancer: applications to the Mexican population. <i>Salud Publica De Mexico</i> , 2011 , 53, 415-9	1.7	1
10	Analytical protocol to identify local ancestry-associated molecular features in cancer. <i>STAR Protocols</i> , 2021 , 2, 100766	1.4	0
9	Differences in somatic TP53 mutation type in breast tumors by race and receptor status.. <i>Breast Cancer Research and Treatment</i> , 2022 , 192, 639	4.4	0
8	Ancestral diversity improves discovery and fine-mapping of genetic loci for anthropometric traits-The Hispanic/Latino Anthropometry Consortium.. <i>Human Genetics and Genomics Advances</i> , 2022 , 3, 100099	0.8	0
7	Response to comment on: Torres-Mejia et al. Moderate-intensity physical activity ameliorates the breast cancer risk in diabetic women. <i>Diabetes Care</i> 2012;35:2500-2502. <i>Diabetes Care</i> , 2013 , 36, e210	14.6	
6	Association between a Polygenic Risk Score for Multiple Myeloma Risk and Overall Survival. <i>Blood</i> , 2019 , 134, 4366-4366	2.2	
5	Germline Variation Predicts Treatment Response in Multiple Myeloma. <i>Blood</i> , 2019 , 134, 4397-4397	2.2	
4	A Genome-Wide Association Study of Myeloma Survival in Utah Uncovers Germline Variants That May Influence Survival of Multiple Myeloma Patients. <i>Blood</i> , 2015 , 126, 2993-2993	2.2	
3	African-American Multiple Myeloma Patients Exhibit Less Bone Disease Compared to Other Racial/Ethnic Groups. <i>Blood</i> , 2010 , 116, 4994-4994	2.2	
2	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	2.2	
1	Genetic determinants of multiple myeloma risk within the Wnt/beta-catenin signaling pathway. <i>Cancer Epidemiology</i> , 2021 , 73, 101972	2.8	