

W Ryan Diver

List of Publications by Citations

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Version: 2024-04-23

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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.

135
papers

12,855
citations

54
h-index

112
g-index

143
ext. papers

14,948
ext. citations

11.9
avg, IF

5.21
L-index

#	Paper	IF	Citations
135	An integrated risk function for estimating the global burden of disease attributable to ambient fine particulate matter exposure. <i>Environmental Health Perspectives</i> , 2014 , 122, 397-403	8.4	1100
134	Global estimates of mortality associated with long-term exposure to outdoor fine particulate matter. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018 , 115, 9592-9597	11.5	810
133	A multistage genome-wide association study in breast cancer identifies two new risk alleles at 1p11.2 and 14q24.1 (RAD51L1). <i>Nature Genetics</i> , 2009 , 41, 579-84	36.3	452
132	A genome-wide association study of lung cancer identifies a region of chromosome 5p15 associated with risk for adenocarcinoma. <i>American Journal of Human Genetics</i> , 2009 , 85, 679-91	11	442
131	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. <i>Nature Genetics</i> , 2013 , 45, 385-91, 391e1-2	36.3	413
130	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. <i>Nature Genetics</i> , 2010 , 42, 978-84	36.3	408
129	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , 2015 , 47, 373-80	36.3	406
128	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. <i>Nature Genetics</i> , 2009 , 41, 585-90	36.3	393
127	Long-Term Ozone Exposure and Mortality in a Large Prospective Study. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2016 , 193, 1134-42	10.2	366
126	Performance of common genetic variants in breast-cancer risk models. <i>New England Journal of Medicine</i> , 2010 , 362, 986-93	59.2	334
125	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. <i>Nature Genetics</i> , 2014 , 46, 1103-9	36.3	331
124	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013 , 45, 392-8, 398e1-2	36.3	327
123	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. <i>Nature Genetics</i> , 2011 , 43, 785-91	36.3	243
122	The landscape of recombination in African Americans. <i>Nature</i> , 2011 , 476, 170-5	50.4	243
121	Relationships between fine particulate air pollution, cardiometabolic disorders, and cardiovascular mortality. <i>Circulation Research</i> , 2015 , 116, 108-15	15.7	241
120	Ischemic Heart Disease Mortality and Long-Term Exposure to Source-Related Components of U.S. Fine Particle Air Pollution. <i>Environmental Health Perspectives</i> , 2016 , 124, 785-94	8.4	223
119	Identification of a new prostate cancer susceptibility locus on chromosome 8q24. <i>Nature Genetics</i> , 2009 , 41, 1055-7	36.3	201

118	Genome-wide association study of renal cell carcinoma identifies two susceptibility loci on 2p21 and 11q13.3. <i>Nature Genetics</i> , 2011 , 43, 60-5	36.3	199
117	Active smoking and breast cancer risk: original cohort data and meta-analysis. <i>Journal of the National Cancer Institute</i> , 2013 , 105, 515-25	9.7	189
116	Genome-wide association study of prostate cancer in men of African ancestry identifies a susceptibility locus at 17q21. <i>Nature Genetics</i> , 2011 , 43, 570-3	36.3	171
115	Dairy, calcium, and vitamin D intake and postmenopausal breast cancer risk in the Cancer Prevention Study II Nutrition Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2005 , 14, 2898-904		156
114	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. <i>Nature Genetics</i> , 2013 , 45, 868-76	36.3	147
113	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
112	Genome-wide association study identifies new prostate cancer susceptibility loci. <i>Human Molecular Genetics</i> , 2011 , 20, 3867-75	5.6	143
111	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, 1252-63	9.7	134
110	A common 8q24 variant in prostate and breast cancer from a large nested case-control study. <i>Cancer Research</i> , 2007 , 67, 2951-6	10.1	127
109	A prospective study of whole grains, fruits, vegetables and colon cancer risk. <i>Cancer Causes and Control</i> , 2003 , 14, 959-70	2.8	123
108	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, 408-15	5.6	109
107	Genome-wide association study identifies multiple susceptibility loci for diffuse large B cell lymphoma. <i>Nature Genetics</i> , 2014 , 46, 1233-8	36.3	108
106	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. <i>Journal of the National Cancer Institute</i> , 2015 , 107, djv279	9.7	107
105	Ambient Air Pollution and Cancer Mortality in the Cancer Prevention Study II. <i>Environmental Health Perspectives</i> , 2017 , 125, 087013	8.4	106
104	Vitamin D pathway gene polymorphisms, diet, and risk of postmenopausal breast cancer: a nested case-control study. <i>Breast Cancer Research</i> , 2007 , 9, R9	8.3	106
103	Genome-wide association study identifies multiple loci associated with bladder cancer risk. <i>Human Molecular Genetics</i> , 2014 , 23, 1387-98	5.6	101
102	Characterizing genetic risk at known prostate cancer susceptibility loci in African Americans. <i>PLoS Genetics</i> , 2011 , 7, e1001387	6	98
101	Outdoor air pollution and cancer: An overview of the current evidence and public health recommendations. <i>Ca-A Cancer Journal for Clinicians</i> , 2020 , 70, 460	220.7	97

100	Comparing the Health Effects of Ambient Particulate Matter Estimated Using Ground-Based versus Remote Sensing Exposure Estimates. <i>Environmental Health Perspectives</i> , 2017 , 125, 552-559	8.4	87
99	Genome-wide meta-analyses of smoking behaviors in African Americans. <i>Translational Psychiatry</i> , 2012 , 2, e119	8.6	86
98	Fine mapping and functional analysis of a common variant in MSMB on chromosome 10q11.2 associated with prostate cancer susceptibility. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009 , 106, 7933-8	11.5	85
97	Common genetic polymorphisms modify the effect of smoking on absolute risk of bladder cancer. <i>Cancer Research</i> , 2013 , 73, 2211-20	10.1	82
96	A genome-wide association study of bladder cancer identifies a new susceptibility locus within SLC14A1, a urea transporter gene on chromosome 18q12.3. <i>Human Molecular Genetics</i> , 2011 , 20, 4282-9	5.6	82
95	A class of non-linear exposure-response models suitable for health impact assessment applicable to large cohort studies of ambient air pollution. <i>Air Quality, Atmosphere and Health</i> , 2016 , 9, 961-972	5.6	79
94	An unusual suspect: an uncommon human-specific synonymous coding variant within the UGT1A6 gene explains a GWAS signal and protects against bladder cancer. <i>Genome Biology</i> , 2011 , 12,	18.3	78
93	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, 6616-33	5.6	77
92	Identification, replication, and fine-mapping of Loci associated with adult height in individuals of african ancestry. <i>PLoS Genetics</i> , 2011 , 7, e1002298	6	77
91	Two susceptibility loci identified for prostate cancer aggressiveness. <i>Nature Communications</i> , 2015 , 6, 6889	17.4	75
90	Genome-wide association study identifies five susceptibility loci for follicular lymphoma outside the HLA region. <i>American Journal of Human Genetics</i> , 2014 , 95, 462-71	11	74
89	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. <i>Human Molecular Genetics</i> , 2012 , 21, 456-62	5.6	74
88	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016 , 7, 10933	17.4	70
87	Common genetic variants in the PSCA gene influence gene expression and bladder cancer risk. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 4974-9	11.5	69
86	Prostate cancer (PCa) risk variants and risk of fatal PCa in the National Cancer Institute Breast and Prostate Cancer Cohort Consortium. <i>European Urology</i> , 2014 , 65, 1069-75	10.2	58
85	Mapping of the UGT1A locus identifies an uncommon coding variant that affects mRNA expression and protects from bladder cancer. <i>Human Molecular Genetics</i> , 2012 , 21, 1918-30	5.6	58
84	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, 2468-75	10.1	58
83	Genome-wide association analysis implicates dysregulation of immunity genes in chronic lymphocytic leukaemia. <i>Nature Communications</i> , 2017 , 8, 14175	17.4	54

82	Work schedule, sleep duration, insomnia, and risk of fatal prostate cancer. <i>American Journal of Preventive Medicine</i> , 2014 , 46, S26-33	6.1	53
81	Generalizability of established prostate cancer risk variants in men of African ancestry. <i>International Journal of Cancer</i> , 2015 , 136, 1210-7	7.5	51
80	Common variation at 2q22.3 (ZEB2) influences the risk of renal cancer. <i>Human Molecular Genetics</i> , 2013 , 22, 825-31	5.6	49
79	Prediction of breast cancer risk by genetic risk factors, overall and by hormone receptor status. <i>Journal of Medical Genetics</i> , 2012 , 49, 601-8	5.8	49
78	Common genetic variants in prostate cancer risk prediction--results from the NCI Breast and Prostate Cancer Cohort Consortium (BPC3). <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012 , 21, 437-44	4	49
77	Characterizing associations and SNP-environment interactions for GWAS-identified prostate cancer risk markers--results from BPC3. <i>PLoS ONE</i> , 2011 , 6, e17142	3.7	49
76	Interactions between cigarette smoking and fine particulate matter in the Risk of Lung Cancer Mortality in Cancer Prevention Study II. <i>American Journal of Epidemiology</i> , 2014 , 180, 1145-9	3.8	48
75	Social Isolation and Mortality in US Black and White Men and Women. <i>American Journal of Epidemiology</i> , 2019 , 188, 102-109	3.8	47
74	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, 437-44	11	46
73	No association between polymorphisms in LEP, LEPR, ADIPOQ, ADIPOR1, or ADIPOR2 and postmenopausal breast cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009 , 18, 2553-7	4	45
72	Use of multivitamins and prostate cancer mortality in a large cohort of US men. <i>Cancer Causes and Control</i> , 2005 , 16, 643-50	2.8	45
71	Interactions between cigarette smoking and ambient PM for cardiovascular mortality. <i>Environmental Research</i> , 2017 , 154, 304-310	7.9	44
70	A genome-wide association study of marginal zone lymphoma shows association to the HLA region. <i>Nature Communications</i> , 2015 , 6, 5751	17.4	44
69	Circadian disruption and fatal ovarian cancer. <i>American Journal of Preventive Medicine</i> , 2014 , 46, S34-41	6.1	43
68	Fine mapping the KLK3 locus on chromosome 19q13.33 associated with prostate cancer susceptibility and PSA levels. <i>Human Genetics</i> , 2011 , 129, 675-85	6.3	41
67	Improved imputation of common and uncommon SNPs with a new reference set. <i>Nature Genetics</i> , 2011 , 44, 6-7	36.3	41
66	Genetically predicted longer telomere length is associated with increased risk of B-cell lymphoma subtypes. <i>Human Molecular Genetics</i> , 2016 , 25, 1663-76	5.6	39
65	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, 3873-84	5.6	39

64	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, 2869-78	5.6	39
63	Identification of novel genetic markers of breast cancer survival. <i>Journal of the National Cancer Institute</i> , 2015 , 107,	9.7	38
62	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. <i>Nature Communications</i> , 2016 , 7, 10979	17.4	37
61	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, 3749-57	5.6	36
60	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, 5603-18	5.6	35
59	Pooled analysis of phosphatidylinositol 3-kinase pathway variants and risk of prostate cancer. <i>Cancer Research</i> , 2010 , 70, 2389-96	10.1	35
58	Genetic variation in candidate obesity genes ADRB2, ADRB3, GHRL, HSD11B1, IRS1, IRS2, and SHC1 and risk for breast cancer in the Cancer Prevention Study II. <i>Breast Cancer Research</i> , 2008 , 10, R57	8.3	35
57	Body mass index, height and risk of lymphoid neoplasms in a large United States cohort. <i>Leukemia and Lymphoma</i> , 2013 , 54, 1221-7	1.9	34
56	The chromosome 2p21 region harbors a complex genetic architecture for association with risk for renal cell carcinoma. <i>Human Molecular Genetics</i> , 2012 , 21, 1190-200	5.6	33
55	Large-scale pathway-based analysis of bladder cancer genome-wide association data from five studies of European background. <i>PLoS ONE</i> , 2012 , 7, e29396	3.7	33
54	The American Cancer Society's Cancer Prevention Study 3 (CPS-3): Recruitment, study design, and baseline characteristics. <i>Cancer</i> , 2017 , 123, 2014-2024	6.4	32
53	Risk factors for fatal breast cancer in African-American women and White women in a large US prospective cohort. <i>American Journal of Epidemiology</i> , 2005 , 162, 734-42	3.8	32
52	Insulin-like growth factor pathway genetic polymorphisms, circulating IGF1 and IGFBP3, and prostate cancer survival. <i>Journal of the National Cancer Institute</i> , 2014 , 106, dju085	9.7	31
51	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, 5260-70	5.6	30
50	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, 1018-27	3.8	29
49	Association of breast cancer risk loci with breast cancer survival. <i>International Journal of Cancer</i> , 2015 , 137, 2837-45	7.5	28
48	Weight loss and postmenopausal breast cancer in a prospective cohort of overweight and obese US women. <i>Cancer Causes and Control</i> , 2011 , 22, 573-9	2.8	28
47	Secondhand Smoke Exposure in Childhood and Adulthood in Relation to Adult Mortality Among Never Smokers. <i>American Journal of Preventive Medicine</i> , 2018 , 55, 345-352	6.1	27

46	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017 , 46, 1814-1822	7.8	27
45	Factors associated with oxidative stress and cancer risk in the Breast and Prostate Cancer Cohort Consortium. <i>Free Radical Research</i> , 2014 , 48, 380-6	4	27
44	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, 1316-22	3.8	27
43	Transforming growth factor beta receptor type I and transforming growth factor beta1 polymorphisms are not associated with postmenopausal breast cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006 , 15, 1236-7	4	27
42	Comprehensive analysis of hormone and genetic variation in 36 genes related to steroid hormone metabolism in pre- and postmenopausal women from the breast and prostate cancer cohort consortium (BPC3). <i>Journal of Clinical Endocrinology and Metabolism</i> , 2011 , 96, E360-7	5.6	26
41	Successful genome-wide scan in paired blood and buccal samples. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007 , 16, 1023-5	4	26
40	Blood levels of cadmium and lead in relation to breast cancer risk in three prospective cohorts. <i>International Journal of Cancer</i> , 2019 , 144, 1010-1016	7.5	25
39	Common germline polymorphisms associated with breast cancer-specific survival. <i>Breast Cancer Research</i> , 2015 , 17, 58	8.3	24
38	Recreational physical activity, leisure sitting time and risk of non-Hodgkin lymphoid neoplasms in the American Cancer Society Cancer Prevention Study II Cohort. <i>International Journal of Cancer</i> , 2012 , 131, 1912-20	7.5	24
37	Residential ambient benzene exposure in the United States and subsequent risk of hematologic malignancies. <i>International Journal of Cancer</i> , 2019 , 145, 2647-2660	7.5	23
36	Large-scale fine mapping of the HNF1B locus and prostate cancer risk. <i>Human Molecular Genetics</i> , 2011 , 20, 3322-9	5.6	22
35	Refining the prostate cancer genetic association within the JAZF1 gene on chromosome 7p15.2. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2010 , 19, 1349-55	4	21
34	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, 212-6	4	21
33	Genetic risk variants associated with in situ breast cancer. <i>Breast Cancer Research</i> , 2015 , 17, 82	8.3	20
32	Residential radon exposure and risk of incident hematologic malignancies in the Cancer Prevention Study-II Nutrition Cohort. <i>Environmental Research</i> , 2016 , 148, 46-54	7.9	20
31	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, 1203-14	5.6	20
30	Whole-exome sequencing of over 4100 men of African ancestry and prostate cancer risk. <i>Human Molecular Genetics</i> , 2016 , 25, 371-81	5.6	19
29	The 19q12 bladder cancer GWAS signal: association with cyclin E function and aggressive disease. <i>Cancer Research</i> , 2014 , 74, 5808-18	10.1	19

28	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	7.5	19
27	HLA Class I and II Diversity Contributes to the Etiologic Heterogeneity of Non-Hodgkin Lymphoma Subtypes. <i>Cancer Research</i> , 2018 , 78, 4086-4096	10.1	18
26	Type II diabetes mellitus and the incidence of epithelial ovarian cancer in the cancer prevention study-II nutrition cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2012 , 21, 2000-5	4	18
25	A genome-wide pleiotropy scan for prostate cancer risk. <i>European Urology</i> , 2015 , 67, 649-57	10.2	17
24	The association between cigarette smoking and non-Hodgkin lymphoid neoplasms in a large US cohort study. <i>Cancer Causes and Control</i> , 2012 , 23, 1231-40	2.8	16
23	Analysis of cohort studies with multivariate and partially observed disease classification data. <i>Biometrika</i> , 2010 , 97, 683-698	2	16
22	Alcohol intake and the incidence of non-hodgkin lymphoid neoplasms in the cancer prevention study II nutrition cohort. <i>American Journal of Epidemiology</i> , 2012 , 176, 60-9	3.8	16
21	Artificially and sugar-sweetened carbonated beverage consumption is not associated with risk of lymphoid neoplasms in older men and women. <i>Journal of Nutrition</i> , 2014 , 144, 2041-9	4.1	15
20	Common variation at 1q24.1 (ALDH9A1) is a potential risk factor for renal cancer. <i>PLoS ONE</i> , 2015 , 10, e0122589	3.7	15
19	Insulin-like Growth Factor Pathway Genetic Polymorphisms, Circulating IGF1 and IGFBP3, and Prostate Cancer Survival. <i>Journal of the National Cancer Institute</i> , 2014 , 106,	9.7	14
18	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
17	Exposure to environmental tobacco smoke and risk of non-Hodgkin lymphoma in nonsmoking men and women. <i>American Journal of Epidemiology</i> , 2014 , 179, 987-95	3.8	12
16	Y chromosome haplogroups and prostate cancer in populations of European and Ashkenazi Jewish ancestry. <i>Human Genetics</i> , 2012 , 131, 1173-85	6.3	11
15	Lupus-related single nucleotide polymorphisms and risk of diffuse large B-cell lymphoma. <i>Lupus Science and Medicine</i> , 2017 , 4, e000187	4.6	10
14	A Prospective Cohort Study of Cigarette Prices and Smoking Cessation in Older Smokers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017 , 26, 1071-1077	4	9
13	Aspirin and other nonsteroidal anti-inflammatory drugs and risk of non-hodgkin lymphoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013 , 22, 422-8	4	9
12	No association between the progesterone receptor gene +331G/A polymorphism and breast cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2004 , 13, 1084-5	4	9
11	Two high-risk susceptibility loci at 6p25.3 and 14q32.13 for Waldenström macroglobulinemia. <i>Nature Communications</i> , 2018 , 9, 4182	17.4	8

10	A genome-wide "pleiotropy scan" does not identify new susceptibility loci for estrogen receptor negative breast cancer. <i>PLoS ONE</i> , 2014 , 9, e85955	3.7	7
9	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	11	7
8	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, 677-89	4.2	6
7	Fine mapping of 14q24.1 breast cancer susceptibility locus. <i>Human Genetics</i> , 2012 , 131, 479-90	6.3	5
6	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
5	Breast cancer risk factors by mode of detection among screened women in the Cancer Prevention Study-II. <i>Breast Cancer Research and Treatment</i> , 2021 , 186, 791-805	4.4	3
4	Erythrocyte levels of cadmium and lead and risk of B-cell non-Hodgkin lymphoma and multiple myeloma. <i>International Journal of Cancer</i> , 2020 , 147, 3110-3118	7.5	1
3	Multilevel-analysis identify a cis-expression quantitative trait locus associated with risk of renal cell carcinoma. <i>Oncotarget</i> , 2015 , 6, 4097-109	3.3	1
2	Frequency of Pathogenic Germline Variants in Cancer-Susceptibility Genes in the Childhood Cancer Survivor Study. <i>JNCI Cancer Spectrum</i> , 2021 , 5, pkab007	4.6	1
1	Association between Smoking Cannabis and Quitting Cigarettes in a Large American Cancer Society Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021 , 30, 1956-1964	4	0