

W Ryan Diver

List of Publications by Year in descending order

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Version: 2024-02-01

141
papers

16,649
citations

27035

58
h-index

17891

125
g-index

143
all docs

143
docs citations

143
times ranked

24958
citing authors

#	ARTICLE	IF	CITATIONS
1	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.	2.8	1,423
2	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.	3.3	1,407
3	Long-Term Ozone Exposure and Mortality in a Large Prospective Study. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2016, 193, 1134-1142.	2.5	602
4	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
5	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. <i>Nature Genetics</i> , 2010, 42, 978-984.	9.4	493
6	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
7	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, 679-691.	2.6	489
8	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, 579-584.	9.4	487
9	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. <i>Nature Genetics</i> , 2009, 41, 585-590.	9.4	434
10	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
11	Performance of Common Genetic Variants in Breast-Cancer Risk Models. <i>New England Journal of Medicine</i> , 2010, 362, 986-993.	13.9	376
12	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , 2013, 45, 392-398.	9.4	374
13	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, 460-479.	157.7	348
14	Relationships Between Fine Particulate Air Pollution, Cardiometabolic Disorders, and Cardiovascular Mortality. <i>Circulation Research</i> , 2015, 116, 108-115.	2.0	327
15	The landscape of recombination in African Americans. <i>Nature</i> , 2011, 476, 170-175.	13.7	319
16	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, 785-794.	2.8	309
17	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
18	Active Smoking and Breast Cancer Risk: Original Cohort Data and Meta-Analysis. <i>Journal of the National Cancer Institute</i> , 2013, 105, 515-525.	3.0	224

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19	Genome-wide association study of renal cell carcinoma identifies two susceptibility loci on 2p21 and 11q13.3. <i>Nature Genetics</i> , 2011, 43, 60-65.	9.4	220
20	Carcinogenicity of night shift work. <i>Lancet Oncology</i> , The, 2019, 20, 1058-1059.	5.1	219
21	Identification of a new prostate cancer susceptibility locus on chromosome 8q24. <i>Nature Genetics</i> , 2009, 41, 1055-1057.	9.4	218
22	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
23	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, 2898-2904.	1.1	188
24	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. <i>Nature Genetics</i> , 2013, 45, 868-876.	9.4	179
25	Ambient Air Pollution and Cancer Mortality in the Cancer Prevention Study II. <i>Environmental Health Perspectives</i> , 2017, 125, 087013.	2.8	169
26	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
27	Genome-wide association study identifies new prostate cancer susceptibility loci. <i>Human Molecular Genetics</i> , 2011, 20, 3867-3875.	1.4	160
28	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.	3.0	152
29	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
30	Genome-wide association study identifies multiple susceptibility loci for diffuse large B cell lymphoma. <i>Nature Genetics</i> , 2014, 46, 1233-1238.	9.4	147
31	A prospective study of whole grains, fruits, vegetables and colon cancer risk. <i>Cancer Causes and Control</i> , 2003, 14, 959-970.	0.8	143
32	Genome-wide association study identifies multiple loci associated with bladder cancer risk. <i>Human Molecular Genetics</i> , 2014, 23, 1387-1398.	1.4	137
33	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
34	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.	2.2	121
35	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
36	Characterizing Genetic Risk at Known Prostate Cancer Susceptibility Loci in African Americans. <i>PLoS Genetics</i> , 2011, 7, e1001387.	1.5	117

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37	Common Genetic Polymorphisms Modify the Effect of Smoking on Absolute Risk of Bladder Cancer. <i>Cancer Research</i> , 2013, 73, 2211-2220.	0.4	107
38	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.	2.8	107
39	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.	1.5	106
40	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, 4282-4289.	1.4	100
41	Fine mapping and functional analysis of a common variant in <i>MSMB</i> 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, 7933-7938.	3.3	96
42	Genome-wide Association Study Identifies Five Susceptibility Loci for Follicular Lymphoma outside the HLA Region. <i>American Journal of Human Genetics</i> , 2014, 95, 462-471.	2.6	96
43	Genome-wide meta-analyses of smoking behaviors in African Americans. <i>Translational Psychiatry</i> , 2012, 2, e119-e119.	2.4	94
44	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016, 7, 10933.	5.8	94
45	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
46	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
47	Two susceptibility loci identified for prostate cancer aggressiveness. <i>Nature Communications</i> , 2015, 6, 6889.	5.8	88
48	Social Isolation and Mortality in US Black and White Men and Women. <i>American Journal of Epidemiology</i> , 2019, 188, 102-109.	1.6	87
49	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. <i>Human Molecular Genetics</i> , 2012, 21, 456-462.	1.4	81
50	Common genetic variants in the <i>PSCA</i> 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, 4974-4979.	3.3	79
51	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, 1069-1075.	0.9	75
52	Genome-wide association analysis implicates dysregulation of immunity genes in chronic lymphocytic leukaemia. <i>Nature Communications</i> , 2017, 8, 14175.	5.8	75
53	Work Schedule, Sleep Duration, Insomnia, and Risk of Fatal Prostate Cancer. <i>American Journal of Preventive Medicine</i> , 2014, 46, S26-S33.	1.6	73
54	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, 1918-1930.	1.4	71

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55	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
56	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
57	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
58	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, 1145-1149.	1.6	61
59	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
60	A genome-wide association study of marginal zone lymphoma shows association to the HLA region. <i>Nature Communications</i> , 2015, 6, 5751.	5.8	58
61	Interactions between cigarette smoking and ambient PM 2.5 for cardiovascular mortality. <i>Environmental Research</i> , 2017, 154, 304-310.	3.7	58
62	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
63	Identification of Novel Genetic Markers of Breast Cancer Survival. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	3.0	56
64	Common variation at 2q22.3 (ZEB2) influences the risk of renal cancer. <i>Human Molecular Genetics</i> , 2013, 22, 825-831.	1.4	54
65	Circadian Disruption and Fatal Ovarian Cancer. <i>American Journal of Preventive Medicine</i> , 2014, 46, S34-S41.	1.6	53
66	Use of multivitamins and prostate cancer mortality in a large cohort of US men. <i>Cancer Causes and Control</i> , 2005, 16, 643-650.	0.8	52
67	Genetically predicted longer telomere length is associated with increased risk of B-cell lymphoma subtypes. <i>Human Molecular Genetics</i> , 2016, 25, 1663-1676.	1.4	52
68	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, 437-444.	1.1	51
69	Fine mapping the KLK3 locus on chromosome 19q13.33 associated with prostate cancer susceptibility and PSA levels. <i>Human Genetics</i> , 2011, 129, 675-685.	1.8	50
70	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
71	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
72	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.	2.2	48

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73	No Association between Polymorphisms in <i>LEP</i> , <i>LEPR</i> , <i>ADIPOQ</i> , <i>ADIPOR1</i> , or <i>ADIPOR2</i> and Postmenopausal Breast Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 2553-2557.	1.1	48
74	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.	1.6	48
75	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
76	Improved imputation of common and uncommon SNPs with a new reference set. <i>Nature Genetics</i> , 2012, 44, 6-7.	9.4	45
77	Body mass index and breast cancer survival: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2017, 46, 1814-1822.	0.9	45
78	Pooled Analysis of Phosphatidylinositol 3-Kinase Pathway Variants and Risk of Prostate Cancer. <i>Cancer Research</i> , 2010, 70, 2389-2396.	0.4	43
79	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
80	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.	2.3	43
81	The American Cancer Society's Cancer Prevention Study 3 (CPS-3): Recruitment, study design, and baseline characteristics. <i>Cancer</i> , 2017, 123, 2014-2024.	2.0	42
82	Body mass index, height and risk of lymphoid neoplasms in a large United States cohort. <i>Leukemia and Lymphoma</i> , 2013, 54, 1221-1227.	0.6	41
83	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, 734-742.	1.6	39
84	Factors associated with oxidative stress and cancer risk in the Breast and Prostate Cancer Cohort Consortium. <i>Free Radical Research</i> , 2014, 48, 380-386.	1.5	38
85	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
86	Successful Genome-Wide Scan in Paired Blood and Buccal Samples. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 1023-1025.	1.1	37
87	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
88	The chromosome 2p21 region harbors a complex genetic architecture for association with risk for renal cell carcinoma. <i>Human Molecular Genetics</i> , 2012, 21, 1190-1200.	1.4	37
89	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
90	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

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91	Residential ambient benzene exposure in the United States and subsequent risk of hematologic malignancies. <i>International Journal of Cancer</i> , 2019, 145, 2647-2660.	2.3	36
92	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.	1.1	36
93	HLA Class I and II Diversity Contributes to the Etiologic Heterogeneity of Non-Hodgkin Lymphoma Subtypes. <i>Cancer Research</i> , 2018, 78, 4086-4096.	0.4	34
94	Weight loss and postmenopausal breast cancer in a prospective cohort of overweight and obese US women. <i>Cancer Causes and Control</i> , 2011, 22, 573-579.	0.8	33
95	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.	3.0	33
96	Association of breast cancer risk <i>loci</i> with breast cancer survival. <i>International Journal of Cancer</i> , 2015, 137, 2837-2845.	2.3	33
97	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
98	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-E367.	1.8	29
99	Transforming Growth Factor Å Receptor Type I and Transforming Growth Factor Å1 Polymorphisms Are Not Associated with Postmenopausal Breast Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 1236-1237.	1.1	28
100	Large-scale fine mapping of the HNF1B locus and prostate cancer risk. <i>Human Molecular Genetics</i> , 2011, 20, 3322-3329.	1.4	28
101	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
102	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
103	Common germline polymorphisms associated with breast cancer-specific survival. <i>Breast Cancer Research</i> , 2015, 17, 58.	2.2	26
104	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.	3.7	26
105	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
106	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, 1912-1920.	2.3	25
107	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, 2041-2049.	1.3	25
108	Genetic risk variants associated with in situ breast cancer. <i>Breast Cancer Research</i> , 2015, 17, 82.	2.2	25

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109	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
110	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
111	Analysis of cohort studies with multivariate and partially observed disease classification data. <i>Biometrika</i> , 2010, 97, 683-698.	1.3	22
112	A Genome-wide Pleiotropy Scan for Prostate Cancer Risk. <i>European Urology</i> , 2015, 67, 649-657.	0.9	21
113	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, 60-69.	1.6	20
114	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, 2000-2005.	1.1	19
115	Common Variation at 1q24.1 (ALDH9A1) Is a Potential Risk Factor for Renal Cancer. <i>PLoS ONE</i> , 2015, 10, e0122589.	1.1	19
116	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
117	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
118	The association between cigarette smoking and non-Hodgkin lymphoid neoplasms in a large US cohort study. <i>Cancer Causes and Control</i> , 2012, 23, 1231-1240.	0.8	17
119	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, .	3.0	16
120	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
121	Lupus-related single nucleotide polymorphisms and risk of diffuse large B-cell lymphoma. <i>Lupus Science and Medicine</i> , 2017, 4, e000187.	1.1	15
122	Two high-risk susceptibility loci at 6p25.3 and 14q32.13 for Waldenström macroglobulinemia. <i>Nature Communications</i> , 2018, 9, 4182.	5.8	15
123	Y chromosome haplogroups and prostate cancer in populations of European and Ashkenazi Jewish ancestry. <i>Human Genetics</i> , 2012, 131, 1173-1185.	1.8	14
124	Exposure to Environmental Tobacco Smoke and Risk of Non-Hodgkin Lymphoma in Nonsmoking Men and Women. <i>American Journal of Epidemiology</i> , 2014, 179, 987-995.	1.6	12
125	Frequency of Pathogenic Germline Variants in Cancer-Susceptibility Genes in the Childhood Cancer Survivor Study. <i>JNCI Cancer Spectrum</i> , 2021, 5, pkab007.	1.4	11
126	A Prospective Cohort Study of Cigarette Prices and Smoking Cessation in Older Smokers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1071-1077.	1.1	10

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127	Aspirin and Other Nonsteroidal Anti-Inflammatory Drugs and Risk of Non-Hodgkin Lymphoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 422-428.	1.1	9
128	No association between the progesterone receptor gene +331G/A polymorphism and breast cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2004, 13, 1084-5.	1.1	9
129	A Genome-Wide <i>â€œ</i> Pleiotropy Scan <i>â€™</i> Does Not Identify New Susceptibility Loci for Estrogen Receptor Negative Breast Cancer. <i>PLoS ONE</i> , 2014, 9, e85955.	1.1	8
130	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.	1.1	8
131	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
132	Erythrocyte levels of cadmium and lead and risk of <i>â€œ</i> cell non <i>â€™</i> Hodgkin lymphoma and multiple myeloma. <i>International Journal of Cancer</i> , 2020, 147, 3110-3118.	2.3	6
133	Fine mapping of 14q24.1 breast cancer susceptibility locus. <i>Human Genetics</i> , 2012, 131, 479-490.	1.8	5
134	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.	1.1	2
135	COVID-19 and Tweets About Quitting Cigarette Smoking: Topic Model Analysis of Twitter Posts 2018-2020. <i>JMIR Infodemiology</i> , 2022, 2, e36215.	1.0	2
136	Multilevel-analysis identify a cis-expression quantitative trait locus associated with risk of renal cell carcinoma. <i>Oncotarget</i> , 2015, 6, 4097-4109.	0.8	1
137	Comparing Remote Sensing, Atmospheric Chemistry and Ground-based Estimates of Fine Particulate Matter on Survival. <i>ISEE Conference Abstracts</i> , 2013, 2013, 5878.	0.0	1
138	Long-Term Ozone Exposure and Cardiovascular Mortality in a Large-Scale Prospective Study. <i>ISEE Conference Abstracts</i> , 2014, 2014, 1679.	0.0	1
139	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, .	3.8	0
140	Improved Imputation of Common and Uncommon Single Nucleotide Polymorphisms (SNPs) with a New Reference Set. <i>Nature Precedings</i> , 2011, , .	0.1	0
141	Analysis of Multivariate Disease Classification Data in the Presence of Partially Missing Disease Traits. <i>Journal of Biometrics & Biostatistics</i> , 2014, 05, .	4.0	0