W Ryan Diver

List of Publications by Year
in descending order

141 papers	16,649 citations	23567 58 h-index	15732 125 g-index
143 all docs	143 docs citations	143 times ranked	22563 citing authors

#	Article	IF	Citations
1	COVID-19 and Tweets About Quitting Cigarette Smoking: Topic Model Analysis of Twitter Posts 2018-2020. JMIR Infodemiology, 2022, 2, e36215.	2.4	2
2	Frequency of Pathogenic Germline Variants in Cancer-Susceptibility Genes in the Childhood Cancer Survivor Study. JNCI Cancer Spectrum, 2021, 5, pkab007.	2.9	11
3	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. American Journal of Human Genetics, 2021, 108, 564-582.	6.2	18
4	Association between Smoking Cannabis and Quitting Cigarettes in a Large American Cancer Society Cohort. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 1956-1964.	2.5	2
5	Breast cancer risk factors by mode of detection among screened women in the Cancer Prevention Study-II. Breast Cancer Research and Treatment, 2021, 186, 791-805.	2.5	8
6	Outdoor air pollution and cancer: An overview of the current evidence and public health recommendations. Ca-A Cancer Journal for Clinicians, 2020, 70, 460-479.	329.8	348
7	Erythrocyte levels of cadmium and lead and risk of <scp>B</scp>â€cell nonâ€Hodgkin lymphoma and multiple myeloma. International Journal of Cancer, 2020, 147, 3110-3118.	5.1	6
8	A meta-analysis of genome-wide association studies of multiple myeloma among men and women of African ancestry. Blood Advances, 2020, 4, 181-190.	5.2	16
9	Blood levels of cadmium and lead in relation to breast cancer risk in three prospective cohorts. International Journal of Cancer, 2019, 144, 1010-1016.	5.1	43
10	Carcinogenicity of night shift work. Lancet Oncology, The, 2019, 20, 1058-1059.	10.7	219
11	Residential ambient benzene exposure in the United States and subsequent risk of hematologic malignancies. International Journal of Cancer, 2019, 145, 2647-2660.	5.1	36
12	Social Isolation and Mortality in US Black and White Men and Women. American Journal of Epidemiology, 2019, 188, 102-109.	3.4	87
13	Two high-risk susceptibility loci at 6p25.3 and 14q32.13 for WaldenstrÃ¶m macroglobulinemia. Nature Communications, 2018, 9, 4182.	12.8	15
14	Global estimates of mortality associated with long-term exposure to outdoor fine particulate matter. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 9592-9597.	7.1	1,407
15	HLA Class I and II Diversity Contributes to the Etiologic Heterogeneity of Non-Hodgkin Lymphoma Subtypes. Cancer Research, 2018, 78, 4086-4096.	0.9	34
16	Secondhand Smoke Exposure in Childhood and Adulthood in Relation to Adult Mortality Among Never Smokers. American Journal of Preventive Medicine, 2018, 55, 345-352.	3.0	48
17	Interactions between cigarette smoking and ambient PM 2.5 for cardiovascular mortality. Environmental Research, 2017, 154, 304-310.	7.5	58
18	The American Cancer Society's Cancer Prevention Study 3 (CPSâ€3): Recruitment, study design, and baseline characteristics. Cancer, 2017, 123, 2014-2024.	4.1	42

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19	Genome-wide association analysis implicates dysregulation of immunity genes in chronic lymphocytic leukaemia. Nature Communications, 2017, 8, 14175.	12.8	75
20	A Prospective Cohort Study of Cigarette Prices and Smoking Cessation in Older Smokers. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 1071-1077.	2.5	10
21	Lupus-related single nucleotide polymorphisms and risk of diffuse large B-cell lymphoma. Lupus Science and Medicine, 2017, 4, e000187.	2.7	15
22	Body mass index and breast cancer survival: a Mendelian randomization analysis. International Journal of Epidemiology, 2017, 46, 1814-1822.	1.9	45
23	Ambient Air Pollution and Cancer Mortality in the Cancer Prevention Study II. Environmental Health Perspectives, 2017, 125, 087013.	6.0	169
24	Comparing the Health Effects of Ambient Particulate Matter Estimated Using Ground-Based versus Remote Sensing Exposure Estimates. Environmental Health Perspectives, 2017, 125, 552-559.	6.0	107
25	Ischemic Heart Disease Mortality and Long-Term Exposure to Source-Related Components of U.S. Fine Particle Air Pollution. Environmental Health Perspectives, 2016, 124, 785-794.	6.0	309
26	A class of non-linear exposure-response models suitable for health impact assessment applicable to large cohort studies of ambient air pollution. Air Quality, Atmosphere and Health, 2016, 9, 961-972.	3.3	106
27	A Meta-analysis of Multiple Myeloma Risk Regions in African and European Ancestry Populations Identifies Putatively Functional Loci. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 1609-1618.	2.5	18
28	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. Nature Communications, 2016, 7, 10933.	12.8	94
29	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. Nature Communications, 2016, 7, 10979.	12.8	50
30	Genetically predicted longer telomere length is associated with increased risk of B-cell lymphoma subtypes. Human Molecular Genetics, 2016, 25, 1663-1676.	2.9	52
31	Residential radon exposure and risk of incident hematologic malignancies in the Cancer Prevention Study-II Nutrition Cohort. Environmental Research, 2016, 148, 46-54.	7.5	26
32	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. Human Molecular Genetics, 2016, 25, 1203-1214.	2.9	38
33	Long-Term Ozone Exposure and Mortality in a Large Prospective Study. American Journal of Respiratory and Critical Care Medicine, 2016, 193, 1134-1142.	5.6	602
34	Whole-exome sequencing of over 4100 men of African ancestry and prostate cancer risk. Human Molecular Genetics, 2016, 25, 371-381.	2.9	26
35	Common germline polymorphisms associated with breast cancer-specific survival. Breast Cancer Research, 2015, 17, 58.	5.0	26
36	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. Journal of the National Cancer Institute, 2015, 107, djv279.	6.3	152

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37	Multilevel-analysis identify a cis-expression quantitative trait locus associated with risk of renal cell carcinoma. Oncotarget, 2015, 6, 4097-4109.	1.8	1
38	A genome-wide association study of marginal zone lymphoma shows association to the HLA region. Nature Communications, 2015, 6, 5751.	12.8	58
39	Association of breast cancer risk loci with breast cancer survival. International Journal of Cancer, 2015, 137, 2837-2845.	5.1	33
40	A Genome-wide Pleiotropy Scan for Prostate Cancer Risk. European Urology, 2015, 67, 649-657.	1.9	21
41	Generalizability of established prostate cancer risk variants in men of <scp>A</scp>frican ancestry. International Journal of Cancer, 2015, 136, 1210-1217.	5.1	62
42	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. Nature Genetics, 2015, 47, 373-380.	21.4	513
43	Genetic risk variants associated with in situ breast cancer. Breast Cancer Research, 2015, 17, 82.	5.0	25
44	Integration of multiethnic fine-mapping and genomic annotation to prioritize candidate functional SNPs at prostate cancer susceptibility regions. Human Molecular Genetics, 2015, 24, 5603-5618.	2.9	50
45	Identification of Novel Genetic Markers of Breast Cancer Survival. Journal of the National Cancer Institute, 2015, 107, .	6.3	56
46	Two susceptibility loci identified for prostate cancer aggressiveness. Nature Communications, 2015, 6, 6889.	12.8	88
47	Relationships Between Fine Particulate Air Pollution, Cardiometabolic Disorders, and Cardiovascular Mortality. Circulation Research, 2015, 116, 108-115.	4.5	327
48	Common Variation at 1q24.1 (ALDH9A1) Is a Potential Risk Factor for Renal Cancer. PLoS ONE, 2015, 10, e0122589.	2.5	19
49	A Genome-Wide â€œPleiotropy Scanâ€•Does Not Identify New Susceptibility Loci for Estrogen Receptor Negative Breast Cancer. PLoS ONE, 2014, 9, e85955.	2.5	8
50	Analysis of Multivariate Disease Classification Data in the Presence of Partially Missing Disease Traits. Journal of Biometrics & Biostatistics, 2014, 05, .	4.0	0
51	Factors associated with oxidative stress and cancer risk in the Breast and Prostate Cancer Cohort Consortium. Free Radical Research, 2014, 48, 380-386.	3.3	38
52	An Integrated Risk Function for Estimating the Global Burden of Disease Attributable to Ambient Fine Particulate Matter Exposure. Environmental Health Perspectives, 2014, 122, 397-403.	6.0	1,423
53	Insulin-like Growth Factor Pathway Genetic Polymorphisms, Circulating IGF1 and IGFBP3, and Prostate Cancer Survival. Journal of the National Cancer Institute, 2014, 106, dju085.	6.3	33
54	Interactions Between Cigarette Smoking and Fine Particulate Matter in the Risk of Lung Cancer Mortality in Cancer Prevention Study II. American Journal of Epidemiology, 2014, 180, 1145-1149.	3.4	61

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55	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. American Journal of Epidemiology, 2014, 180, 1018-1027.	3.4	36
56	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. Human Molecular Genetics, 2014, 23, 6616-6633.	2.9	90
57	Post-GWAS geneâ€“environment interplay in breast cancer: results from the Breast and Prostate Cancer Cohort Consortium and a meta-analysis on 79 000 women. Human Molecular Genetics, 2014, 23, 5260-5270.	2.9	37
58	Artificially and Sugar-Sweetened Carbonated Beverage Consumption Is Not Associated with Risk of Lymphoid Neoplasms in Older Men and Women. Journal of Nutrition, 2014, 144, 2041-2049.	2.9	25
59	Insulin-like Growth Factor Pathway Genetic Polymorphisms, Circulating IGF1 and IGFBP3, and Prostate Cancer Survival. Journal of the National Cancer Institute, 2014, 106, .	6.3	16
60	Genome-wide association study identifies multiple loci associated with bladder cancer risk. Human Molecular Genetics, 2014, 23, 1387-1398.	2.9	137
61	Exposure to Environmental Tobacco Smoke and Risk of Non-Hodgkin Lymphoma in Nonsmoking Men and Women. American Journal of Epidemiology, 2014, 179, 987-995.	3.4	12
62	Prostate Cancer (PCa) Risk Variants and Risk of Fatal PCa in the National Cancer Institute Breast and Prostate Cancer Cohort Consortium. European Urology, 2014, 65, 1069-1075.	1.9	75
63	The 19q12 Bladder Cancer GWAS Signal: Association with Cyclin E Function and Aggressive Disease. Cancer Research, 2014, 74, 5808-5818.	0.9	24
64	Genome-wide association study identifies multiple susceptibility loci for diffuse large B cell lymphoma. Nature Genetics, 2014, 46, 1233-1238.	21.4	147
65	Genome-wide Association Study Identifies Five Susceptibility Loci for Follicular Lymphoma outside the HLA Region. American Journal of Human Genetics, 2014, 95, 462-471.	6.2	96
66	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. Nature Genetics, 2014, 46, 1103-1109.	21.4	408
67	Genome-wide Scan of 29,141 African Americans Finds No Evidence of Directional Selection since Admixture. American Journal of Human Genetics, 2014, 95, 437-444.	6.2	69
68	Circadian Disruption and Fatal Ovarian Cancer. American Journal of Preventive Medicine, 2014, 46, S34-S41.	3.0	53
69	Work Schedule, Sleep Duration, Insomnia, and Risk of Fatal Prostate Cancer. American Journal of Preventive Medicine, 2014, 46, S26-S33.	3.0	73
70	Long-Term Ozone Exposure and Cardiovascular Mortality in a Large-Scale Prospective Study. ISEE Conference Abstracts, 2014, 2014, 1679.	0.0	1
71	Common Genetic Polymorphisms Modify the Effect of Smoking on Absolute Risk of Bladder Cancer. Cancer Research, 2013, 73, 2211-2220.	0.9	107
72	Active Smoking and Breast Cancer Risk: Original Cohort Data and Meta-Analysis. Journal of the National Cancer Institute, 2013, 105, 515-525.	6.3	224

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73	One thousand genomes imputation in the national cancer institute breast and prostate cancer cohort consortium aggressive prostate cancer genomeâ€wide association study. Prostate, 2013, 73, 677-689.	2.3	6
74	Insulinâ€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). International Journal of Cancer, 2013, 133, 495-504.	5.1	28
75	Genome-wide association studies identify four ER negativeâ€“specific breast cancer risk loci. Nature Genetics, 2013, 45, 392-398.	21.4	374
76	Identification of 23 new prostate cancer susceptibility loci using the iCOGS custom genotyping array. Nature Genetics, 2013, 45, 385-391.	21.4	492
77	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. Nature Genetics, 2013, 45, 868-876.	21.4	179
78	A meta-analysis of genome-wide association studies to identify prostate cancer susceptibility loci associated with aggressive and non-aggressive disease. Human Molecular Genetics, 2013, 22, 408-415.	2.9	118
79	Common variation at 2q22.3 (ZEB2) influences the risk of renal cancer. Human Molecular Genetics, 2013, 22, 825-831.	2.9	54
80	Aspirin and Other Nonsteroidal Anti-Inflammatory Drugs and Risk of Non-Hodgkin Lymphoma. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 422-428.	2.5	9
81	Body mass index, height and risk of lymphoid neoplasms in a large United States cohort. Leukemia and Lymphoma, 2013, 54, 1221-1227.	1.3	41
82	Comparing Remote Sensing, Atmospheric Chemistry and Ground-based Estimates of Fine Particulate Matter on Survival. ISEE Conference Abstracts, 2013, 2013, 5878.	0.0	1
83	Mapping of the UGT1A locus identifies an uncommon coding variant that affects mRNA expression and protects from bladder cancer. Human Molecular Genetics, 2012, 21, 1918-1930.	2.9	71
84	Type II Diabetes Mellitus and the Incidence of Epithelial Ovarian Cancer in the Cancer Prevention Study-II Nutrition Cohort. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 2000-2005.	2.5	19
85	Replication of Five Prostate Cancer Loci Identified in an Asian Populationâ€”Results from the NCI Breast and Prostate Cancer Cohort Consortium (BPC3). Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 212-216.	2.5	23
86	Prediction of breast cancer risk by genetic risk factors, overall and by hormone receptor status. Journal of Medical Genetics, 2012, 49, 601-608.	3.2	58
87	Improved imputation of common and uncommon SNPs with a new reference set. Nature Genetics, 2012, 44, 6-7.	21.4	45
88	Genome-wide meta-analyses of smoking behaviors in African Americans. Translational Psychiatry, 2012, 2, e119-e119.	4.8	94
89	Common genetic variants in the PSCA gene influence gene expression and bladder cancer risk. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 4974-4979.	7.1	79
90	Common Genetic Variants in Prostate Cancer Risk Predictionâ€”Results from the NCI Breast and Prostate Cancer Cohort Consortium (BPC3). Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 437-444.	2.5	51

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91	Alcohol Intake and the Incidence of Non-Hodgkin Lymphoid Neoplasms in the Cancer Prevention Study II Nutrition Cohort. American Journal of Epidemiology, 2012, 176, 60-69.	3.4	20
92	The chromosome 2p21 region harbors a complex genetic architecture for association with risk for renal cell carcinoma. Human Molecular Genetics, 2012, 21, 1190-1200.	2.9	37
93	A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. Human Molecular Genetics, 2012, 21, 5373-5384.	2.9	168
94	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. Human Molecular Genetics, 2012, 21, 456-462.	2.9	81
95	Recreational physical activity, leisure sitting time and risk of nonâ€Hodgkin lymphoid neoplasms in the American Cancer Society Cancer Prevention Study II Cohort. International Journal of Cancer, 2012, 131, 1912-1920.	5.1	25
96	The association between cigarette smoking and non-Hodgkin lymphoid neoplasms in a large US cohort study. Cancer Causes and Control, 2012, 23, 1231-1240.	1.8	17
97	Y chromosome haplogroups and prostate cancer in populations of European and Ashkenazi Jewish ancestry. Human Genetics, 2012, 131, 1173-1185.	3.8	14
98	Fine mapping of 14q24.1 breast cancer susceptibility locus. Human Genetics, 2012, 131, 479-490.	3.8	5
99	Large-Scale Pathway-Based Analysis of Bladder Cancer Genome-Wide Association Data from Five Studies of European Background. PLoS ONE, 2012, 7, e29396.	2.5	36
100	Interactions Between Genetic Variants and Breast Cancer Risk Factors in the Breast and Prostate Cancer Cohort Consortium. Journal of the National Cancer Institute, 2011, 103, 1252-1263.	6.3	147
101	N-Acetyltransferase 2 Polymorphisms, Tobacco Smoking, and Breast Cancer Risk in the Breast and Prostate Cancer Cohort Consortium. American Journal of Epidemiology, 2011, 174, 1316-1322.	3.4	31
102	An unusual suspect: an uncommon human-specific synonymous coding variant within the UGT1A6 gene explains a GWAS signal and protects against bladder cancer. Genome Biology, 2011, 12, .	8.8	0
103	Improved Imputation of Common and Uncommon Single Nucleotide Polymorphisms (SNPs) with a New Reference Set. Nature Precedings, 2011, , .	0.1	0
104	Genome-wide association study of renal cell carcinoma identifies two susceptibility loci on 2p21 and 11q13.3. Nature Genetics, 2011, 43, 60-65.	21.4	220
105	Genome-wide association study of prostate cancer in men of African ancestry identifies a susceptibility locus at 17q21. Nature Genetics, 2011, 43, 570-573.	21.4	198
106	Weight loss and postmenopausal breast cancer in a prospective cohort of overweight and obese US women. Cancer Causes and Control, 2011, 22, 573-579.	1.8	33
107	Fine mapping the KLK3 locus on chromosome 19q13.33 associated with prostate cancer susceptibility and PSA levels. Human Genetics, 2011, 129, 675-685.	3.8	50
108	Seven prostate cancer susceptibility loci identified by a multi-stage genome-wide association study. Nature Genetics, 2011, 43, 785-791.	21.4	265

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109	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). Journal of Clinical Endocrinology and Metabolism, 2011, 96, E360-E367.	3.6	29
110	Genome-wide association study identifies new prostate cancer susceptibility loci. Human Molecular Genetics, 2011, 20, 3867-3875.	2.9	160
111	Fine mapping of a region of chromosome 11q13 reveals multiple independent loci associated with risk of prostate cancer. Human Molecular Genetics, 2011, 20, 2869-2878.	2.9	43
112	Large-scale fine mapping of the HNF1B locus and prostate cancer risk. Human Molecular Genetics, 2011, 20, 3322-3329.	2.9	28
113	The landscape of recombination in African Americans. Nature, 2011, 476, 170-175.	27.8	319
114	A genome-wide association study of bladder cancer identifies a new susceptibility locus within SLC14A1, a urea transporter gene on chromosome 18q12.3. Human Molecular Genetics, 2011, 20, 4282-4289.	2.9	100
115	Characterizing Genetic Risk at Known Prostate Cancer Susceptibility Loci in African Americans. PLoS Genetics, 2011, 7, e1001387.	3.5	117
116	Identification, Replication, and Fine-Mapping of Loci Associated with Adult Height in Individuals of African Ancestry. PLoS Genetics, 2011, 7, e1002298.	3.5	93
117	Characterizing Associations and SNP-Environment Interactions for GWAS-Identified Prostate Cancer Risk Markersâ€"Results from BPC3. PLoS ONE, 2011, 6, e17142.	2.5	57
118	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. Nature Genetics, 2010, 42, 978-984.	21.4	493
119	Pooled Analysis of Phosphatidylinositol 3-Kinase Pathway Variants and Risk of Prostate Cancer. Cancer Research, 2010, 70, 2389-2396.	0.9	43
120	Refining the Prostate Cancer Genetic Association within the JAZF1 Gene on Chromosome 7p15.2. Cancer Epidemiology Biomarkers and Prevention, 2010, 19, 1349-1355.	2.5	26
121	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 consortiumâ€. Human Molecular Genetics, 2010, 19, 3873-3884.	2.9	45
122	Analysis of cohort studies with multivariate and partially observed disease classification data. Biometrika, 2010, 97, 683-698.	2.4	22
123	Performance of Common Genetic Variants in Breast-Cancer Risk Models. New England Journal of Medicine, 2010, 362, 986-993.	27.0	376
124	No Association between Polymorphisms in LEP, LEPR, ADIPOQ, ADIPOR1, or ADIPOR2 and Postmenopausal Breast Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 2553-2557.	2.5	48
125	Fine mapping and functional analysis of a common variant in MSMB on chromosome 10q11.2 associated with prostate cancer susceptibility. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 7933-7938.	7.1	96
126	Quantitative trait loci predicting circulating sex steroid hormones in men from the NCI-Breast and Prostate Cancer Cohort Consortium (BPC3). Human Molecular Genetics, 2009, 18, 3749-3757.	2.9	37

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127	A multistage genome-wide association study in breast cancer identifies two new risk alleles at 1p11.2 and 14q24.1 (RAD51L1). Nature Genetics, 2009, 41, 579-584.	21.4	487
128	Newly discovered breast cancer susceptibility loci on 3p24 and 17q23.2. Nature Genetics, 2009, 41, 585-590.	21.4	434
129	Identification of a new prostate cancer susceptibility locus on chromosome 8q24. Nature Genetics, 2009, 41, 1055-1057.	21.4	218
130	A Genome-wide Association Study of Lung Cancer Identifies a Region of Chromosome 5p15 Associated with Risk for Adenocarcinoma. American Journal of Human Genetics, 2009, 85, 679-691.	6.2	489
131	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. Breast Cancer Research, 2008, 10, R57.	5.0	48
132	A Common 8q24 Variant in Prostate and Breast Cancer from a Large Nested Case-Control Study. Cancer Research, 2007, 67, 2951-2956.	0.9	136
133	Vitamin D pathway gene polymorphisms, diet, and risk of postmenopausal breast cancer: a nested case-control study. Breast Cancer Research, 2007, 9, R9.	5.0	121
134	Successful Genome-Wide Scan in Paired Blood and Buccal Samples. Cancer Epidemiology Biomarkers and Prevention, 2007, 16, 1023-1025.	2.5	37
135	Haplotype Analysis of the HSD17B1 Gene and Risk of Breast Cancer: A Comprehensive Approach to Multicenter Analyses of Prospective Cohort Studies. Cancer Research, 2006, 66, 2468-2475.	0.9	64
136	Transforming Growth Factor Â Receptor Type I and Transforming Growth Factor Â1 Polymorphisms Are Not Associated with Postmenopausal Breast Cancer. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 1236-1237.	2.5	28
137	Use of multivitamins and prostate cancer mortality in a large cohort of US men. Cancer Causes and Control, 2005, 16, 643-650.	1.8	52
138	Risk Factors for Fatal Breast Cancer in African-American Women and White Women in a Large US Prospective Cohort. American Journal of Epidemiology, 2005, 162, 734-742.	3.4	39
139	Dairy, Calcium, and Vitamin D Intake and Postmenopausal Breast Cancer Risk in the Cancer Prevention Study II Nutrition Cohort. Cancer Epidemiology Biomarkers and Prevention, 2005, 14, 2898-2904.	2.5	188
140	No association between the progesterone receptor gene +331G/A polymorphism and breast cancer. Cancer Epidemiology Biomarkers and Prevention, 2004, 13, 1084-5.	2.5	9
141	A prospective study of whole grains, fruits, vegetables and colon cancer risk. Cancer Causes and Control, 2003, 14, 959-970.	1.8	143