

Daniel O Stram

List of Publications by Year in descending order

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Version: 2024-02-01

167
papers

11,973
citations

44069

48
h-index

31849

101
g-index

167
all docs

167
docs citations

167
times ranked

19462
citing authors

#	ARTICLE	IF	CITATIONS
1	Prognostic utility of self-reported sarcopenia (SARC-F) in the Multiethnic Cohort. <i>Journal of Cachexia, Sarcopenia and Muscle</i> , 2022, 13, 987-1002.	7.3	8
2	Outdoor ambient air pollution and breast cancer survival among California participants of the Multiethnic Cohort Study. <i>Environment International</i> , 2022, 161, 107088.	10.0	8
3	Predicted gene expression in ancestrally diverse populations leads to discovery of susceptibility loci for lifestyle and cardiometabolic traits. <i>American Journal of Human Genetics</i> , 2022, 109, 669-679.	6.2	5
4	Lung Cancer in the Mayak Workers Cohort: Risk Estimation and Uncertainty Analysis. <i>Radiation Research</i> , 2021, 195, 334-346.	1.5	17
5	Associations of the gut microbiome with hepatic adiposity in the Multiethnic Cohort Adiposity Phenotype Study. <i>Gut Microbes</i> , 2021, 13, 1965463.	9.8	16
6	Ethnic Differences of Urinary Cadmium in Cigarette Smokers from the Multiethnic Cohort Study. <i>International Journal of Environmental Research and Public Health</i> , 2021, 18, 2669.	2.6	1
7	Ethnic differences in excretion of butadiene-DNA adducts by current smokers. <i>Carcinogenesis</i> , 2021, 42, 694-704.	2.8	6
8	Urinary phthalate exposures and risk of breast cancer: the Multiethnic Cohort study. <i>Breast Cancer Research</i> , 2021, 23, 44.	5.0	33
9	Genetic discovery and risk characterization in type 2 diabetes across diverse populations. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100029.	1.7	23
10	Racial/ethnic disparities in weight or BMI change in adulthood and pancreatic cancer incidence: The multiethnic cohort. <i>Cancer Medicine</i> , 2021, 10, 4097-4106.	2.8	4
11	Risk of breast cancer and prediagnostic urinary excretion of bisphenol A, triclosan and parabens: The Multiethnic Cohort Study. <i>International Journal of Cancer</i> , 2021, 149, 1426-1434.	5.1	21
12	Genome-Wide Association Analyses Identify Variants in IRF4 Associated With Acute Myeloid Leukemia and Myelodysplastic Syndrome Susceptibility. <i>Frontiers in Genetics</i> , 2021, 12, 554948.	2.3	8
13	Association between Airport-Related Ultrafine Particles and Risk of Malignant Brain Cancer: A Multiethnic Cohort Study. <i>Cancer Research</i> , 2021, 81, 4360-4369.	0.9	5
14	Genome-wide association study of pancreatic fat: The Multiethnic Cohort Adiposity Phenotype Study. <i>PLoS ONE</i> , 2021, 16, e0249615.	2.5	2
15	Multiethnic Prediction of Nicotine Biomarkers and Association With Nicotine Dependence. <i>Nicotine and Tobacco Research</i> , 2021, 23, 2162-2169.	2.6	6
16	Discovery of structural deletions in breast cancer predisposition genes using whole genome sequencing data from 2000 women of African-ancestry. <i>Human Genetics</i> , 2021, 140, 1449-1457.	3.8	4
17	Novel genetic variants associated with mortality after unrelated donor allogeneic hematopoietic cell transplantation. <i>EClinicalMedicine</i> , 2021, 40, 101093.	7.1	8
18	The association between ambient air pollutants and pancreatic cancer in the Multiethnic Cohort Study. <i>Environmental Research</i> , 2021, 202, 111608.	7.5	8

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19	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 623-642.	2.5	19
20	Association between ambient air pollution and breast cancer risk: The multiethnic cohort study. <i>International Journal of Cancer</i> , 2020, 146, 699-711.	5.1	60
21	Urinary N7-(1-hydroxy-3-buten-2-yl) guanine adducts in humans: temporal stability and association with smoking. <i>Mutagenesis</i> , 2020, 35, 19-26.	2.6	13
22	Issues in Interpreting Epidemiologic Studies of Populations Exposed to Low-Dose, High-Energy Photon Radiation. <i>Journal of the National Cancer Institute Monographs</i> , 2020, 2020, 176-187.	2.1	27
23	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. <i>Nature Communications</i> , 2020, 11, 3833.	12.8	88
24	A case-control study of the joint effect of reproductive factors and radiation treatment for first breast cancer and risk of contralateral breast cancer in the WECARE study. <i>Breast</i> , 2020, 54, 62-69.	2.2	3
25	Estrogen Plus Progestin Hormone Therapy and Ovarian Cancer. <i>Epidemiology</i> , 2020, 31, 402-408.	2.7	12
26	Exposure measurement error in air pollution studies: the impact of shared, multiplicative measurement error on epidemiological health risk estimates. <i>Air Quality, Atmosphere and Health</i> , 2020, 13, 631-643.	3.3	7
27	Association Between Outdoor Air Pollution and Risk of Malignant and Benign Brain Tumors: The Multiethnic Cohort Study. <i>JNCI Cancer Spectrum</i> , 2020, 4, pkz107.	2.9	16
28	Minority-centric meta-analyses of blood lipid levels identify novel loci in the Population Architecture using Genomics and Epidemiology (PAGE) study. <i>PLoS Genetics</i> , 2020, 16, e1008684.	3.5	17
29	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.	5.2	16
30	Meta-Analysis of Genome-Wide Association Studies of Acute Myeloid Leukemia (AML) Patients Identifies Variants Associated with Risk of 11q23/KMT2A-Translocated and Core-Binding Factor (CBF) AML and Suggests a Role for Transcription Elongation in Leukemogenesis. <i>Blood</i> , 2020, 136, 29-30.	1.4	0
31	Characteristics of and Risk Factors for Monoclonal Gammopathy of Undetermined Significance (MGUS) in the Multiethnic Cohort Study. <i>Blood</i> , 2020, 136, 28-29.	1.4	0
32	Title is missing!. , 2020, 16, e1008684.		0
33	Title is missing!. , 2020, 16, e1008684.		0
34	Title is missing!. , 2020, 16, e1008684.		0
35	Title is missing!. , 2020, 16, e1008684.		0
36	Title is missing!. , 2020, 16, e1008684.		0

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37	Title is missing!. , 2020, 16, e1008684.		0
38	Pancreatic Cancer Following Incident Diabetes in African Americans and Latinos: The Multiethnic Cohort. Journal of the National Cancer Institute, 2019, 111, 27-33.	6.3	51
39	Association of a Pathway-Specific Genetic Risk Score With Risk of Radiation-Associated Contralateral Breast Cancer. JAMA Network Open, 2019, 2, e1912259.	5.9	5
40	Association between mitochondrial genetic variation and breast cancer risk: The Multiethnic Cohort. PLoS ONE, 2019, 14, e0222284.	2.5	6
41	Genetic analyses of diverse populations improves discovery for complex traits. Nature, 2019, 570, 514-518.	27.8	679
42	Interethnic differences in pancreatic cancer incidence and risk factors: The Multiethnic Cohort. Cancer Medicine, 2019, 8, 3592-3603.	2.8	35
43	Propensity for Intra-abdominal and Hepatic Adiposity Varies Among Ethnic Groups. Gastroenterology, 2019, 156, 966-975.e10.	1.3	80
44	Exposure measurement error in air pollution studies: A framework for assessing shared, multiplicative measurement error in ensemble learning estimates of nitrogen oxides. Environment International, 2019, 125, 97-106.	10.0	11
45	Validation of genetic associations with acute GVHD and nonrelapse mortality in DISCOVeRY-BMT. Blood Advances, 2019, 3, 2337-2341.	5.2	8
46	Multiple functional variants in the IL1RL1 region are pretransplant markers for risk of GVHD and infection deaths. Blood Advances, 2019, 3, 2512-2524.	5.2	7
47	Exome Chip Meta-analysis Fine Maps Causal Variants and Elucidates the Genetic Architecture of Rare Coding Variants in Smoking and Alcohol Use. Biological Psychiatry, 2019, 85, 946-955.	1.3	69
48	De Novo and Therapy-Related Acute Myeloid Leukemia and Myelodysplastic Syndrome: Similarities and Differences in SNP-Array Detected Chromosomal Aberrations in Pre-Transplant Blood Samples. Blood, 2019, 134, 1430-1430.	1.4	2
49	Comprehensive Investigation of White Blood Cell and Gene Expression Profiles As Risk Factors for Multiple Myeloma in African Americans. Blood, 2019, 134, 4379-4379.	1.4	0
50	Genome Wide Interaction Analysis Identifies Expression Quantitative Trait Loci Associated with Reduced Survival after Reduced Intensity Conditioning HLA-Matched Unrelated Donor Allogeneic Hematopoietic Cell Transplant. Blood, 2019, 134, 4595-4595.	1.4	0
51	Exome chip analyses identify genes affecting mortality after HLA-matched unrelated-donor blood and marrow transplantation. Blood, 2018, 131, 2490-2499.	1.4	21
52	Breast Cancer Family History and Contralateral Breast Cancer Risk in Young Women: An Update From the Women's Environmental Cancer and Radiation Epidemiology Study. Journal of Clinical Oncology, 2018, 36, 1513-1520.	1.6	44
53	Growth factor genes and change in mammographic density after stopping combined hormone therapy in the California Teachers Study. BMC Cancer, 2018, 18, 1072.	2.6	1
54	Evaluation of 71 Coronary Artery Disease Risk Variants in a Multiethnic Cohort. Frontiers in Cardiovascular Medicine, 2018, 5, 19.	2.4	13

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55	Association of internal smoking dose with blood DNA methylation in three racial/ethnic populations. <i>Clinical Epigenetics</i> , 2018, 10, 110.	4.1	34
56	Genome Wide Association Analyses Identify Pleiotropic Variants Associated with Acute Myeloid Leukemia (AML) and Myelodysplastic Syndrome (MDS) Susceptibility. <i>Blood</i> , 2018, 132, 1500-1500.	1.4	0
57	Multiple Functional Donor Polymorphisms in IL1RL1 region Associate with Death Due to GvHD or Infection after Unrelated Donor Allogeneic Hematopoietic Stem Cell Transplantation (HCT) for AML and MDS. <i>Blood</i> , 2018, 132, 312-312.	1.4	0
58	Coffee Drinking and Alcoholic and Nonalcoholic Fatty Liver Diseases and Viral Hepatitis in the Multiethnic Cohort. <i>Clinical Gastroenterology and Hepatology</i> , 2017, 15, 1305-1307.	4.4	22
59	<i>BRCA2</i> Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , 2017, 77, 2789-2799.	0.9	75
60	Genetic Determinants of 1,3-Butadiene Metabolism and Detoxification in Three Populations of Smokers with Different Risks of Lung Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1034-1042.	2.5	22
61	Characterizing Genetic Susceptibility to Breast Cancer in Women of African Ancestry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1016-1026.	2.5	24
62	Novel colon cancer susceptibility variants identified from a genome-wide association study in African Americans. <i>International Journal of Cancer</i> , 2017, 140, 2728-2733.	5.1	26
63	Isotope Dilution nanoLC/ESI-MS/MS Quantitation of Urinary N7-(1-Hydroxy-3-buten-2-yl) Guanine Adducts in Humans and Their Use as Biomarkers of Exposure to 1,3-Butadiene. <i>Chemical Research in Toxicology</i> , 2017, 30, 678-688.	3.3	21
64	Multi-SNP Haplotype Analysis Methods for Association Analysis. <i>Methods in Molecular Biology</i> , 2017, 1666, 485-504.	0.9	19
65	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017, 551, 92-94.	27.8	1,099
66	Replication and validation of genetic polymorphisms associated with survival after allogeneic blood or marrow transplant. <i>Blood</i> , 2017, 130, 1585-1596.	1.4	45
67	THE GROWTH OF BIOSTATISTICS AND ESTIMATION OF CANCER RISK ESTIMATES: PAST, CURRENT, AND FUTURE CHALLENGES. <i>Radiation Protection Dosimetry</i> , 2017, 173, 32-35.	0.8	0
68	Mortality from solid cancers other than lung, liver, and bone in relation to external dose among plutonium and non-plutonium workers in the Mayak Worker Cohort. <i>Radiation and Environmental Biophysics</i> , 2017, 56, 121-125.	1.4	16
69	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	6.3	57
70	Genetic association with B-cell acute lymphoblastic leukemia in allogeneic transplant patients differs by age and sex. <i>Blood Advances</i> , 2017, 1, 1717-1728.	5.2	15
71	Impact of common genetic determinants of Hemoglobin A1c on type 2 diabetes risk and diagnosis in ancestrally diverse populations: A transethnic genome-wide meta-analysis. <i>PLoS Medicine</i> , 2017, 14, e1002383.	8.4	341
72	Correction of confidence intervals in excess relative risk models using Monte Carlo dosimetry systems with shared errors. <i>PLoS ONE</i> , 2017, 12, e0174641.	2.5	32

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73	Sex and ethnic/racial-specific risk factors for gallbladder disease. BMC Gastroenterology, 2017, 17, 153.	2.0	64
74	Association of CYP2A6 activity with lung cancer incidence in smokers: The multiethnic cohort study. PLoS ONE, 2017, 12, e0178435.	2.5	35
75	Metabolites of the Polycyclic Aromatic Hydrocarbon Phenanthrene in the Urine of Cigarette Smokers from Five Ethnic Groups with Differing Risks for Lung Cancer. PLoS ONE, 2016, 11, e0156203.	2.5	23
76	Prevalence of chronic liver disease and cirrhosis by underlying cause in understudied ethnic groups: The multiethnic cohort. Hepatology, 2016, 64, 1969-1977.	7.3	237
77	Replication of associations between genetic polymorphisms and chronic graft-versus-host disease. Blood, 2016, 128, 2450-2456.	1.4	32
78	No Association between the Mitochondrial Genome and Prostate Cancer Risk: The Multiethnic Cohort. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 1001-1003.	2.5	15
79	Breast Cancer Risk From Modifiable and Nonmodifiable Risk Factors Among White Women in the United States. JAMA Oncology, 2016, 2, 1295.	7.1	285
80	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
81	Novel Association of Genetic Markers Affecting CYP2A6 Activity and Lung Cancer Risk. Cancer Research, 2016, 76, 5768-5776.	0.9	57
82	Meta-analysis of Rare Variant Association Tests in Multiethnic Populations. Genetic Epidemiology, 2016, 40, 57-65.	1.3	9
83	Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. Cancer Discovery, 2016, 6, 1052-1067.	9.4	157
84	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. Nature Communications, 2016, 7, 11843.	12.8	86
85	Fine scale mapping of the 17q22 breast cancer locus using dense SNPs, genotyped within the Collaborative Oncological Gene-Environment Study (COGs). Scientific Reports, 2016, 6, 32512.	3.3	19
86	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. Nature Communications, 2016, 7, 10979.	12.8	50
87	The contribution of rare variation to prostate cancer heritability. Nature Genetics, 2016, 48, 30-35.	21.4	139
88	Prostate Cancer Susceptibility in Men of African Ancestry at 8q24. Journal of the National Cancer Institute, 2016, 108, djv431.	6.3	111
89	Whole-exome sequencing of over 4100 men of African ancestry and prostate cancer risk. Human Molecular Genetics, 2016, 25, 371-381.	2.9	26
90	HLA Haplotypes Are Associated with Multiple Myeloma Risk in the African American Multiple Myeloma Study (AAMMS). Blood, 2016, 128, 3250-3250.	1.4	1

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91	Benzene Uptake and Glutathione S-transferase T1 Status as Determinants of S-Phenylmercapturic Acid in Cigarette Smokers in the Multiethnic Cohort. PLoS ONE, 2016, 11, e0150641.	2.5	20
92	Replication of Candidate SNP Survival Analyses and Gene-Based Tests of Association with Survival Outcomes after an Unrelated Donor Blood or Marrow Transplant: Results from the Discovery-BMT Study. Blood, 2016, 128, 71-71.	1.4	0
93	Exome Array Analyses Identify New Genes Influencing Survival Outcomes after HLA-Matched Unrelated Donor Blood and Marrow Transplantation. Blood, 2016, 128, 518-518.	1.4	0
94	Exome Array Analyses Identify Low-Frequency Germline Variants Associated with Increased Risk of AML in a HLA-Matched Unrelated Donor Blood and Marrow Transplant Population. Blood, 2016, 128, 42-42.	1.4	0
95	ABO blood group alleles and prostate cancer risk: Results from the breast and prostate cancer cohort consortium (BPC3). Prostate, 2015, 75, 1677-1681.	2.3	14
96	Methodological Considerations in Estimation of Phenotype Heritability Using Genome-Wide SNP Data, Illustrated by an Analysis of the Heritability of Height in a Large Sample of African Ancestry Adults. PLoS ONE, 2015, 10, e0131106.	2.5	2
97	Association of Genes, Pathways, and Haplogroups of the Mitochondrial Genome with the Risk of Colorectal Cancer: The Multiethnic Cohort. PLoS ONE, 2015, 10, e0136796.	2.5	27
98	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. American Journal of Human Genetics, 2015, 96, 487-497.	6.2	101
99	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. Human Molecular Genetics, 2015, 24, 2966-2984.	2.9	40
100	The Contribution of Common Genetic Variation to Nicotine and Cotinine Glucuronidation in Multiple Ethnic/Racial Populations. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 119-127.	2.5	47
101	Generalizability of established prostate cancer risk variants in men of African ancestry. International Journal of Cancer, 2015, 136, 1210-1217.	5.1	62
102	Vitamin D-Associated Genetic Variation and Risk of Breast Cancer in the Breast and Prostate Cancer Cohort Consortium (BPC3). Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 627-630.	2.5	20
103	Fine-Scale Mapping of the 5q11.2 Breast Cancer Locus Reveals at Least Three Independent Risk Variants Regulating MAP3K1. American Journal of Human Genetics, 2015, 96, 5-20.	6.2	76
104	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
105	Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. Clinical Cancer Research, 2015, 21, 5264-5276.	7.0	33
106	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
107	Evaluating the ovarian cancer gonadotropin hypothesis: A candidate gene study. Gynecologic Oncology, 2015, 136, 542-548.	1.4	15
108	Variation in Levels of the Lung Carcinogen NNAL and Its Glucuronides in the Urine of Cigarette Smokers from Five Ethnic Groups with Differing Risks for Lung Cancer. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 561-569.	2.5	39

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109	Trans-ancestry genome-wide association study identifies 12 genetic loci influencing blood pressure and implicates a role for DNA methylation. <i>Nature Genetics</i> , 2015, 47, 1282-1293.	21.4	294
110	Pleiotropic Analysis of Cancer Risk Loci on Esophageal Adenocarcinoma Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1801-1803.	2.5	7
111	Population Distribution of Lifetime Risk of Ovarian Cancer in the United States. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 671-676.	2.5	82
112	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. <i>Human Molecular Genetics</i> , 2015, 24, 285-298.	2.9	38
113	Shared Dosimetry Error in Epidemiological Dose-Response Analyses. <i>PLoS ONE</i> , 2015, 10, e0119418.	2.5	34
114	Mercapturic Acids Derived from the Toxicants Acrolein and Crotonaldehyde in the Urine of Cigarette Smokers from Five Ethnic Groups with Differing Risks for Lung Cancer. <i>PLoS ONE</i> , 2015, 10, e0124841.	2.5	56
115	Combined Donor and Recipient Non-HLA Genotypes Show Evidence of Genome Wide Association with Transplant Related Mortality (TRM) after HLA-Matched Unrelated Donor Blood and Marrow Transplantation (URD-BMT) (DISCOVeRY-BMT study). <i>Blood</i> , 2015, 126, 61-61.	1.4	7
116	Evidence for Heterogeneous Genetic Associations with Acute Lymphoblastic Leukemia (ALL) By Cytogenetics and Sex in High-Risk Patients Treated with Matched Unrelated Donor Allogeneic Blood or Marrow Transplant (URD-BMT). <i>Blood</i> , 2015, 126, 2621-2621.	1.4	5
117	Genome-Wide Association Study of Overall and Progression-Free Survival after HLA-Matched Unrelated Donor Blood and Marrow Transplantation (DISCOVeRY-BMT study). <i>Blood</i> , 2015, 126, 397-397.	1.4	1
118	Pleiotropy of Cancer Susceptibility Variants on the Risk of Non-Hodgkin Lymphoma: The PAGE Consortium. <i>PLoS ONE</i> , 2014, 9, e89791.	2.5	16
119	Multiple Nonglycemic Genomic Loci Are Newly Associated With Blood Level of Glycated Hemoglobin in East Asians. <i>Diabetes</i> , 2014, 63, 2551-2562.	0.6	61
120	Nicotine N-glucuronidation relative to N-oxidation and C-oxidation and UGT2B10 genotype in five ethnic/racial groups. <i>Carcinogenesis</i> , 2014, 35, 2526-2533.	2.8	124
121	Analysis of Cancer Risks in Populations Near Nuclear Facilities. <i>Health Physics</i> , 2014, 106, 305-306.	0.5	4
122	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014, 23, 6616-6633.	2.9	90
123	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , 2014, 23, 6096-6111.	2.9	53
124	Trans-ethnic genome-wide association study of colorectal cancer identifies a new susceptibility locus in VTIIA. <i>Nature Communications</i> , 2014, 5, 4613.	12.8	72
125	A comprehensive examination of breast cancer risk loci in African American women. <i>Human Molecular Genetics</i> , 2014, 23, 5518-5526.	2.9	42
126	Diabetes and Racial/Ethnic Differences in Hepatocellular Carcinoma Risk: The Multiethnic Cohort. <i>Journal of the National Cancer Institute</i> , 2014, 106, dju326-dju326.	6.3	44

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127	Leveraging population admixture to characterize the heritability of complex traits. <i>Nature Genetics</i> , 2014, 46, 1356-1362.	21.4	69
128	High Calorie Intake Is Associated With Worsening Insulin Resistance and β -Cell Function in Hispanic Women After Gestational Diabetes Mellitus. <i>Diabetes Care</i> , 2014, 37, 3294-3300.	8.6	26
129	The Impact of GWAS Findings on Cancer Etiology and Prevention. <i>Current Epidemiology Reports</i> , 2014, 1, 130-137.	2.4	3
130	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. <i>Nature Genetics</i> , 2014, 46, 1103-1109.	21.4	408
131	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , 2014, 5, 4999.	12.8	105
132	Fine-Mapping α -IGF1 and Prostate Cancer Risk in African Americans: The Multiethnic Cohort Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 1928-1932.	2.5	7
133	Joint Effects of Known Type 2 Diabetes Susceptibility Loci in Genome-Wide Association Study of Singapore Chinese: The Singapore Chinese Health Study. <i>PLoS ONE</i> , 2014, 9, e87762.	2.5	15
134	Genetic Susceptibility Markers of Multiple Myeloma in African-Americans. <i>Blood</i> , 2014, 124, 2030-2030.	1.4	0
135	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , 2013, 45, 371-384.	21.4	493
136	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , 2013, 45, 353-361.	21.4	960
137	Multi-SNP Haplotype Analysis Methods for Association Analysis. <i>Methods in Molecular Biology</i> , 2012, 850, 423-452.	0.9	17
138	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , 2012, 44, 651-658.	21.4	519
139	Improved Imputation of Common and Uncommon Single Nucleotide Polymorphisms (SNPs) with a New Reference Set. <i>Nature Precedings</i> , 2011, , .	0.1	0
140	Age-specific effects of hormone therapy use on overall mortality and ischemic heart disease mortality among women in the California Teachers Study. <i>Menopause</i> , 2011, 18, 253-261.	2.0	24
141	Characterizing Genetic Risk at Known Prostate Cancer Susceptibility Loci in African Americans. <i>PLoS Genetics</i> , 2011, 7, e1001387.	3.5	117
142	Beta-Cryptoxanthin and Lung Cancer in Shanghai, China—An Examination of Potential Confounding with Cigarette Smoking Using Urinary Cotinine as a Biomarker for True Tobacco Exposure. <i>Nutrition and Cancer</i> , 2007, 57, 123-129.	2.0	7
143	Prostate cancer incidence and intake of fruits, vegetables and related micronutrients: the multiethnic cohort study* (United States). <i>Cancer Causes and Control</i> , 2006, 17, 1193-1207.	1.8	68
144	Software for tag single nucleotide polymorphism selection. <i>Human Genomics</i> , 2005, 2, 144.	2.9	35

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145	Genetic variation in the HSD17B1 gene and risk of prostate cancer. PLoS Genetics, 2005, preprint, e68.	3.5	6
146	Designs for Studies of Personal Exposure to Air Pollution and the Impact of Measurement Error. Journal of Toxicology and Environmental Health - Part A: Current Issues, 2005, 68, 1181-1187.	2.3	2
147	Tag SNP selection for association studies. Genetic Epidemiology, 2004, 27, 365-374.	1.3	165
148	Power and Uncertainty Analysis of Epidemiological Studies of Radiation-Related Disease Risk in which Dose Estimates are Based on a Complex Dosimetry System: Some Observations. Radiation Research, 2003, 160, 408-417.	1.5	64
149	Modeling and E-M Estimation of Haplotype-Specific Relative Risks from Genotype Data for a Case-Control Study of Unrelated Individuals. Human Heredity, 2003, 55, 179-190.	0.8	249
150	Choosing Haplotype-Tagging SNPS Based on Unphased Genotype Data Using a Preliminary Sample of Unrelated Subjects with an Example from the Multiethnic Cohort Study. Human Heredity, 2003, 55, 27-36.	0.8	386
151	Is Residual Confounding a Reasonable Explanation for the Apparent Protective Effects of Beta-carotene Found in Epidemiologic Studies of Lung Cancer in Smokers?. American Journal of Epidemiology, 2002, 155, 622-628.	3.4	62
152	Bone Marrow Leukemic Progenitor Cell Content in Pediatric T-Lineage Acute Lymphoblastic Leukemia Patients with an Isolated Extramedullar First Relapse. Leukemia and Lymphoma, 2001, 40, 279-285.	1.3	4
153	Opsoclonus-myoclonus-ataxia syndrome in neuroblastoma: Clinical outcome and antineuronal antibodies? a report from the children's cancer group study. Medical and Pediatric Oncology, 2001, 36, 612-622.	1.0	203
154	International neuroblastoma pathology classification for prognostic evaluation of patients with peripheral neuroblastic tumors. Cancer, 2001, 92, 2451-2461.	4.1	273
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