Daniel O Stram

List of Publications by Citations

165 papers	8,766 citations	42 h-index	92 g-index
167 ext. papers	10,661 ext. citations	7.5 avg, IF	5.15 L-index

#	Paper	IF	Citations
165	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. *Nature Genetics*, **2013**, 45, 353-61, 361e1-2	36.3	813
164	Association analysis identifies 65 new breast cancer risk loci. *Nature*, **2017**, 551, 92-94	50.4	643
163	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. *Nature Genetics*, **2013**, 45, 371-84, 384e1-2	36.3	422
162	Detectable clonal mosaicism and its relationship to aging and cancer. *Nature Genetics*, **2012**, 44, 651-8	36.3	409
161	Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. *Nature Genetics*, **2015**, 47, 373-80	36.3	406
160	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	1.1	371
159	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. *Nature Genetics*, **2014**, 46, 1103-9	36.3	331
158	Singapore Chinese Health Study: development, validation, and calibration of the quantitative food frequency questionnaire. *Nutrition and Cancer*, **2001**, 39, 187-95	2.8	299
157	Genetic analyses of diverse populations improves discovery for complex traits. *Nature*, **2019**, 570, 514-518	50.4	291
156	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, 179-90	1.1	230
155	Trans-ancestry genome-wide association study identifies 12 genetic loci influencing blood pressure and implicates a role for DNA methylation. *Nature Genetics*, **2015**, 47, 1282-1293	36.3	223
154	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. *PLoS Medicine*, **2017**, 14, e1002383	11.6	223
153	International neuroblastoma pathology classification for prognostic evaluation of patients with peripheral neuroblastic tumors: a report from the Children's Cancer Group. *Cancer*, **2001**, 92, 2451-61	6.4	220
152	Breast Cancer Risk From Modifiable and Nonmodifiable Risk Factors Among White Women in the United States. *JAMA Oncology*, **2016**, 2, 1295-1302	13.4	189
151	Prevalence of chronic liver disease and cirrhosis by underlying cause in understudied ethnic groups: The multiethnic cohort. *Hepatology*, **2016**, 64, 1969-1977	11.2	175
150	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, 612-22		157
149	Tag SNP selection for association studies. *Genetic Epidemiology*, **2004**, 27, 365-74	2.6	149

148	Neuroblastoma in adults and adolescents: an indolent course with poor survival. *Cancer*, **1997**, 79, 2028-35	6.4	118
147	Histopathology (International Neuroblastoma Pathology Classification) and MYCN status in patients with peripheral neuroblastic tumors: a report from the Children's Cancer Group. *Cancer*, **2001**, 92, 2699-708	6.4	110
146	The contribution of rare variation to prostate cancer heritability. *Nature Genetics*, **2016**, 48, 30-5	36.3	106
145	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, 1052-67	24.4	104
144	Nicotine N-glucuronidation relative to N-oxidation and C-oxidation and UGT2B10 genotype in five ethnic/racial groups. *Carcinogenesis*, **2014**, 35, 2526-33	4.6	103
143	Characterizing genetic risk at known prostate cancer susceptibility loci in African Americans. *PLoS Genetics*, **2011**, 7, e1001387	6	98
142	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. *Nature Communications*, **2014**, 4, 4999	17.4	87
141	Characterization of large structural genetic mosaicism in human autosomes. *American Journal of Human Genetics*, **2015**, 96, 487-97	11	77
140	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, 6616-33	5.6	77
139	Prostate Cancer Susceptibility in Men of African Ancestry at 8q24. *Journal of the National Cancer Institute*, **2016**, 108,	9.7	72
138	Population distribution of lifetime risk of ovarian cancer in the United States. *Cancer Epidemiology Biomarkers and Prevention*, **2015**, 24, 671-676	4	67
137	Latency analysis in epidemiologic studies of occupational exposures: application to the Colorado Plateau uranium miners cohort. *American Journal of Industrial Medicine*, **1999**, 35, 246-56	2.7	66
136	Trans-ethnic genome-wide association study of colorectal cancer identifies a new susceptibility locus in VTI1A. *Nature Communications*, **2014**, 5, 4613	17.4	62
135	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. *Nature Communications*, **2016**, 7, 11843	17.4	59
134	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	11	59
133	Prostate cancer incidence and intake of fruits, vegetables and related micronutrients: the multiethnic cohort study* (United States). *Cancer Causes and Control*, **2006**, 17, 1193-207	2.8	57
132	Comprehensive analysis of chromosome 1p deletions in neuroblastoma. *Medical and Pediatric Oncology*, **2001**, 36, 32-6		57
131	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, 408-17	3.1	52

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130	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, 622-8	3.8	52
129	Generalizability of established prostate cancer risk variants in men of African ancestry. *International Journal of Cancer*, **2015**, 136, 1210-7	7.5	51
128	Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. *Cancer Research*, **2017**, 77, 2789-2799	10.1	49
127	Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. *Human Molecular Genetics*, **2014**, 23, 6096-111	5.6	48
126	Multiple nonglycemic genomic loci are newly associated with blood level of glycated hemoglobin in East Asians. *Diabetes*, **2014**, 63, 2551-62	0.9	46
125	Leveraging population admixture to characterize the heritability of complex traits. *Nature Genetics*, **2014**, 46, 1356-62	36.3	45
124	Propensity for Intra-abdominal and Hepatic Adiposity Varies Among Ethnic Groups. *Gastroenterology*, **2019**, 156, 966-975.e10	13.3	44
123	The contribution of common genetic variation to nicotine and cotinine glucuronidation in multiple ethnic/racial populations. *Cancer Epidemiology Biomarkers and Prevention*, **2015**, 24, 119-27	4	42
122	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. *PLoS ONE*, **2015**, 10, e0124841	3.7	42
121	Allelic deletion at chromosome bands 11q14-23 is common in neuroblastoma. *Medical and Pediatric Oncology*, **2001**, 36, 24-7		41
120	Novel Association of Genetic Markers Affecting CYP2A6 Activity and Lung Cancer Risk. *Cancer Research*, **2016**, 76, 5768-5776	10.1	41
119	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. *Journal of the National Cancer Institute*, **2017**, 109,	9.7	38
118	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. *Nature Communications*, **2016**, 7, 10979	17.4	37
117	Replication and validation of genetic polymorphisms associated with survival after allogeneic blood or marrow transplant. *Blood*, **2017**, 130, 1585-1596	2.2	36
116	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. *Human Molecular Genetics*, **2015**, 24, 2966-84	5.6	36
115	Integration of multiethnic fine-mapping and genomic annotation to prioritize candidate functional SNPs at prostate cancer susceptibility regions. *Human Molecular Genetics*, **2015**, 24, 5603-18	5.6	35
114	Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. *Human Molecular Genetics*, **2015**, 24, 285-98	5.6	35
113	Pancreatic Cancer Following Incident Diabetes in African Americans and Latinos: The Multiethnic Cohort. *Journal of the National Cancer Institute*, **2019**, 111, 27-33	9.7	35

112	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	7.9	35
111	Diabetes and racial/ethnic differences in hepatocellular carcinoma risk: the multiethnic cohort. *Journal of the National Cancer Institute*, **2014**, 106,	9.7	34
110	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, 561-9	4	32
109	Sex and ethnic/racial-specific risk factors for gallbladder disease. *BMC Gastroenterology*, **2017**, 17, 153	3	31
108	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. *Nature Communications*, **2020**, 11, 3833	17.4	31
107	Software for tag single nucleotide polymorphism selection. *Human Genomics*, **2005**, 2, 144-51	6.8	30
106	Shared dosimetry error in epidemiological dose-response analyses. *PLoS ONE*, **2015**, 10, e0119418	3.7	30
105	Breast Cancer Family History and Contralateral Breast Cancer Risk in Young Women: An Update From the Women@ Environmental Cancer and Radiation Epidemiology Study. *Journal of Clinical Oncology*, **2018**, 36, 1513-1520	2.2	29
104	A comprehensive examination of breast cancer risk loci in African American women. *Human Molecular Genetics*, **2014**, 23, 5518-26	5.6	28
103	Association of CYP2A6 activity with lung cancer incidence in smokers: The multiethnic cohort study. *PLoS ONE*, **2017**, 12, e0178435	3.7	28
102	Association between ambient air pollution and breast cancer risk: The multiethnic cohort study. *International Journal of Cancer*, **2020**, 146, 699-711	7.5	28
101	Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. *Clinical Cancer Research*, **2015**, 21, 5264-76	12.9	24
100	Correction of confidence intervals in excess relative risk models using Monte Carlo dosimetry systems with shared errors. *PLoS ONE*, **2017**, 12, e0174641	3.7	24
99	Treatment of overt isolated testicular relapse in children on therapy for acute lymphoblastic leukemia. A report from the Childrens Cancer Group. *Cancer*, **1994**, 73, 219-23	6.4	24
98	Replication of associations between genetic polymorphisms and chronic graft-versus-host disease. *Blood*, **2016**, 128, 2450-2456	2.2	24
97	Interethnic differences in pancreatic cancer incidence and risk factors: The Multiethnic Cohort. *Cancer Medicine*, **2019**, 8, 3592-3603	4.8	23
96	Association of internal smoking dose with blood DNA methylation in three racial/ethnic populations. *Clinical Epigenetics*, **2018**, 10, 110	7.7	22
95	Age-specific effects of hormone therapy use on overall mortality and ischemic heart disease mortality among women in the California Teachers Study. *Menopause*, **2011**, 18, 253-61	2.5	21

#	Reference		
94	Association of Genes, Pathways, and Haplogroups of the Mitochondrial Genome with the Risk of Colorectal Cancer: The Multiethnic Cohort. *PLoS ONE*, **2015**, 10, e0136796	3.7	20
93	Whole-exome sequencing of over 4100 men of African ancestry and prostate cancer risk. *Human Molecular Genetics*, **2016**, 25, 371-81	5.6	19
92	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, 627-30	4	18
91	Coffee Drinking and Alcoholic and Nonalcoholic Fatty Liver Diseases and Viral Hepatitis in the Multiethnic Cohort. *Clinical Gastroenterology and Hepatology*, **2017**, 15, 1305-1307	6.9	17
90	Novel colon cancer susceptibility variants identified from a genome-wide association study in African Americans. *International Journal of Cancer*, **2017**, 140, 2728-2733	7.5	17
89	Multidrug resistance-associated protein 1 (MRP1) expression in neuroblastoma cell lines and primary tumors. *Medical and Pediatric Oncology*, **2000**, 35, 619-22		17
88	Issues in Interpreting Epidemiologic Studies of Populations Exposed to Low-Dose, High-Energy Photon Radiation. *Journal of the National Cancer Institute Monographs*, **2020**, 2020, 176-187	4.8	17
87	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	3.7	17
86	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	4.9	16
85	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	3.7	16
84	Genetic Determinants of 1,3-Butadiene Metabolism and Detoxification in Three Populations of Smokers with Different Risks of Lung Cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2017**, 26, 1034-1042	4	15
83	Multi-SNP Haplotype Analysis Methods for Association Analysis. *Methods in Molecular Biology*, **2017**, 1666, 485-504	1.4	15
82	Isotope Dilution nanoLC/ESI-HRMS 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. *Chemical Research in Toxicology*, **2017**, 30, 678-688	4	14
81	High calorie intake is associated with worsening insulin resistance and ß-cell function in Hispanic women after gestational diabetes mellitus. *Diabetes Care*, **2014**, 37, 3294-300	14.6	14
80	Pleiotropy of cancer susceptibility variants on the risk of non-Hodgkin lymphoma: the PAGE consortium. *PLoS ONE*, **2014**, 9, e89791	3.7	14
79	Joint effects of known type 2 diabetes susceptibility loci in genome-wide association study of Singapore Chinese: the Singapore Chinese health study. *PLoS ONE*, **2014**, 9, e87762	3.7	14
78	Exome chip analyses identify genes affecting mortality after HLA-matched unrelated-donor blood and marrow transplantation. *Blood*, **2018**, 131, 2490-2499	2.2	13
77	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	4	13

76	Characterizing Genetic Susceptibility to Breast Cancer in Women of African Ancestry. *Cancer Epidemiology Biomarkers and Prevention*, **2017**, 26, 1016-1026	4	12
75	Evaluating the ovarian cancer gonadotropin hypothesis: a candidate gene study. *Gynecologic Oncology*, **2015**, 136, 542-8	4.9	12
74	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. *Radiation and Environmental Biophysics*, **2017**, 56, 121-125	2	12
73	A strategic view of randomized trial design in low-incidence paediatric cancer. *Statistics in Medicine*, **1999**, 18, 1183-97	2.3	12
72	No Association between the Mitochondrial Genome and Prostate Cancer Risk: The Multiethnic Cohort. *Cancer Epidemiology Biomarkers and Prevention*, **2016**, 25, 1001-3	4	12
71	Association Between Outdoor Air Pollution and Risk of Malignant and Benign Brain Tumors: The Multiethnic Cohort Study. *JNCI Cancer Spectrum*, **2020**, 4, pkz107	4.6	10
70	Genetic association with B-cell acute lymphoblastic leukemia in allogeneic transplant patients differs by age and sex. *Blood Advances*, **2017**, 1, 1717-1728	7.8	10
69	ABO blood group alleles and prostate cancer risk: Results from the breast and prostate cancer cohort consortium (BPC3). *Prostate*, **2015**, 75, 1677-81	4.2	10
68	Urinary N7-(1-hydroxy-3-buten-2-yl) guanine adducts in humans: temporal stability and association with smoking. *Mutagenesis*, **2020**, 35, 19-26	2.8	9
67	Evaluation of 71 Coronary Artery Disease Risk Variants in a Multiethnic Cohort. *Frontiers in Cardiovascular Medicine*, **2018**, 5, 19	5.4	8
66	Lung Cancer in the Mayak Workers Cohort: Risk Estimation and Uncertainty Analysis. *Radiation Research*, **2021**, 195, 334-346	3.1	8
65	Pleiotropic analysis of cancer risk loci on esophageal adenocarcinoma risk. *Cancer Epidemiology Biomarkers and Prevention*, **2015**, 24, 1801-3	4	7
64	Multi-SNP haplotype analysis methods for association analysis. *Methods in Molecular Biology*, **2012**, 850, 423-52	1.4	7
63	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. *Nutrition and Cancer*, **2007**, 57, 123-9	2.8	7
62	Genetic discovery and risk characterization in type 2 diabetes across diverse populations. *Human Genetics and Genomics Advances*, **2021**, 2, 100029-100029	0.8	7
61	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	12.9	6
60	Exposure Measurement Error in Air Pollution Studies: The Impact of Shared, Multiplicative Measurement Error on Epidemiological Health Risk Estimates. *Air Quality, Atmosphere and Health*, **2020**, 13, 631-643	5.6	6
59	Genetic variation in the HSD17B1 gene and risk of prostate cancer. *PLoS Genetics*, **2005**, preprint, e68	6	6

58	Meta-Analysis of Rare Variant Association Tests in Multiethnic Populations. *Genetic Epidemiology*, **2016**, 40, 57-65	2.6	6
57	Minority-centric meta-analyses of blood lipid levels identify novel loci in the Population Architecture using Genomics and Epidemiology (PAGE) study. *PLoS Genetics*, **2020**, 16, e1008684	6	5
56	Fine-mapping IGF1 and prostate cancer risk in African Americans: the multiethnic cohort study. *Cancer Epidemiology Biomarkers and Prevention*, **2014**, 23, 1928-32	4	5
55	A meta-analysis of genome-wide association studies of multiple myeloma among men and women of African ancestry. *Blood Advances*, **2020**, 4, 181-190	7.8	5
54	A generalized estimating equations approach to fitting major gene models in segregation analysis of continuous phenotypes. *Genetic Epidemiology*, **1993**, 10, 61-74	2.6	4
53	Urinary phthalate exposures and risk of breast cancer: the Multiethnic Cohort study. *Breast Cancer Research*, **2021**, 23, 44	8.3	4
52	Validation of genetic associations with acute GVHD and nonrelapse mortality in DISCOVeRY-BMT. *Blood Advances*, **2019**, 3, 2337-2341	7.8	4
51	Multiple functional variants in the region are pretransplant markers for risk of GVHD and infection deaths. *Blood Advances*, **2019**, 3, 2512-2524	7.8	4
50	Breast Cancer Risk Factors and Survival by Tumor Subtype: Pooled Analyses from the Breast Cancer Association Consortium. *Cancer Epidemiology Biomarkers and Prevention*, **2021**, 30, 623-642	4	4
49	The Impact of GWAS Findings on Cancer Etiology and Prevention. *Current Epidemiology Reports*, **2014**, 1, 130-137	2.9	3
48	Analysis of cancer risks in populations near nuclear facilities: Phase I. A report by the National Academies Nuclear and Radiation Studies Board. *Health Physics*, **2014**, 106, 305-6	2.3	3
47	Bone marrow leukemic progenitor cell content in pediatric T-lineage acute lymphoblastic leukemia patients with an isolated extramedullary first relapse. *Leukemia and Lymphoma*, **2001**, 40, 279-85	1.9	3
46	Estrogen Plus Progestin Hormone Therapy and Ovarian Cancer: A Complicated Relationship Explored. *Epidemiology*, **2020**, 31, 402-408	3.1	3
45	Ethnic differences in excretion of butadiene-DNA adducts by current smokers. *Carcinogenesis*, **2021**, 42, 694-704	4.6	3
44	Genome-Wide Association Analyses Identify Variants in Associated With Acute Myeloid Leukemia and Myelodysplastic Syndrome Susceptibility. *Frontiers in Genetics*, **2021**, 12, 554948	4.5	3
43	The association between ambient air pollutants and pancreatic cancer in the Multiethnic Cohort Study. *Environmental Research*, **2021**, 202, 111608	7.9	3
42	International neuroblastoma pathology classification for prognostic evaluation of patients with peripheral neuroblastic tumors **2001**, 92, 2451		3
41	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	3.7	2

40	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, 1181-7	3.2	2
39	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. *Breast*, **2020**, 54, 62-69	3.6	2
38	Racial/ethnic disparities in weight or BMI change in adulthood and pancreatic cancer incidence: The multiethnic cohort. *Cancer Medicine*, **2021**, 10, 4097-4106	4.8	2
37	Association of a Pathway-Specific Genetic Risk Score With Risk of Radiation-Associated Contralateral Breast Cancer. *JAMA Network Open*, **2019**, 2, e1912259	10.4	1
36	Association between mitochondrial genetic variation and breast cancer risk: The Multiethnic Cohort. *PLoS ONE*, **2019**, 14, e0222284	3.7	1
35	Prognostic utility of self-reported sarcopenia (SARC-F) in the Multiethnic Cohort.. *Journal of Cachexia, Sarcopenia and Muscle*, **2022**,	10.3	1
34	Outdoor ambient air pollution and breast cancer survival among California participants of the Multiethnic Cohort Study.. *Environment International*, **2022**, 161, 107088	12.9	1
33	HLA Haplotypes Are Associated with Multiple Myeloma Risk in the African American Multiple Myeloma Study (AAMMS). *Blood*, **2016**, 128, 3250-3250	2.2	1
32	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). *Blood*, **2015**, 126, 61-61	2.2	1
31	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). *Blood*, **2015**, 126, 2621-2621	2.2	1
30	Risk of breast cancer and prediagnostic urinary excretion of bisphenol A, triclosan and parabens: The Multiethnic Cohort Study. *International Journal of Cancer*, **2021**, 149, 1426-1434	7.5	1
29	Associations of the gut microbiome with hepatic adiposity in the Multiethnic Cohort Adiposity Phenotype Study. *Gut Microbes*, **2021**, 13, 1965463	8.8	1
28	Growth factor genes and change in mammographic density after stopping combined hormone therapy in the California Teachers Study. *BMC Cancer*, **2018**, 18, 1072	4.8	1
27	Discovery of structural deletions in breast cancer predisposition genes using whole genome sequencing data from > 2000 women of African-ancestry. *Human Genetics*, **2021**, 140, 1449-1457	6.3	1
26	Histopathology (International Neuroblastoma Pathology Classification) and MYCN status in patients with peripheral neuroblastic tumors **2001**, 92, 2699		1
25	Predicted gene expression in ancestrally diverse populations leads to discovery of susceptibility loci for lifestyle and cardiometabolic traits.. *American Journal of Human Genetics*, **2022**,	11	1
24	Association between Airport-Related Ultrafine Particles and Risk of Malignant Brain Cancer: A Multiethnic Cohort Study. *Cancer Research*, **2021**, 81, 4360-4369	10.1	0
23	Genome-wide association study of pancreatic fat: The Multiethnic Cohort Adiposity Phenotype Study. *PLoS ONE*, **2021**, 16, e0249615	3.7	0

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22	Multiethnic Prediction of Nicotine Biomarkers and Association With Nicotine Dependence. *Nicotine and Tobacco Research*, **2021**, 23, 2162-2169	4.9	0
21	Novel genetic variants associated with mortality after unrelated donor allogeneic hematopoietic cell transplantation. *EClinicalMedicine*, **2021**, 40, 101093	11.3	0
20	THE GROWTH OF BIOSTATISTICS AND ESTIMATION OF CANCER RISK ESTIMATES: PAST, CURRENT, AND FUTURE CHALLENGES. *Radiation Protection Dosimetry*, **2017**, 173, 32-35	0.9	
19	Kolonel and Stram Respond to Riboli and Kaaks. *American Journal of Epidemiology*, **2000**, 151, 375-376	3.8	
18	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. *Blood*, **2020**, 136, 29-30	2.2	
17	Characteristics of and Risk Factors for Monoclonal Gammopathy of Undetermined Significance (MGUS) in the Multiethnic Cohort Study. *Blood*, **2020**, 136, 28-29	2.2	
16	Genome Wide Association Analyses Identify Pleiotropic Variants Associated with Acute Myeloid Leukemia (AML) and Myelodysplastic Syndrome (MDS) Susceptibility. *Blood*, **2018**, 132, 1500-1500	2.2	
15	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. *Blood*, **2018**, 132, 312-312	2.2	
14	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	2.2	
13	Comprehensive Investigation of White Blood Cell and Gene Expression Profiles As Risk Factors for Multiple Myeloma in African Americans. *Blood*, **2019**, 134, 4379-4379	2.2	
12	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	2.2	
11	Genetic Susceptibility Markers of Multiple Myeloma in African-Americans. *Blood*, **2014**, 124, 2030-2030	2.2	
10	Genome-Wide Association Study of Overall and Progression-Free Survival after HLA-Matched Unrelated Donor Blood and Marrow Transplantation (DISCOVeRY-BMT study). *Blood*, **2015**, 126, 397-397	2.2	
9	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	2.2	
8	Exome Array Analyses Identify New Genes Influencing Survival Outcomes after HLA-Matched Unrelated Donor Blood and Marrow Transplantation. *Blood*, **2016**, 128, 518-518	2.2	
7	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	2.2	
6	Minority-centric meta-analyses of blood lipid levels identify novel loci in the Population Architecture using Genomics and Epidemiology (PAGE) study **2020**, 16, e1008684		
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4   Minority-centric meta-analyses of blood lipid levels identify novel loci in the Population
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