Daniel O Stram

List of Publications by Citations

200 papers	19,498 citations	67 h-index	137 g-index
204 ext. papers	22,189 ext. citations	8.7 avg, IF	5.73 L-index

#	Paper	IF	Citations
200	Treatment of high-risk neuroblastoma with intensive chemotherapy, radiotherapy, autologous bone marrow transplantation, and 13-cis-retinoic acid. Children's Cancer Group. *New England Journal of Medicine*, **1999**, 341, 1165-73	59.2	1502
199	Large-scale genotyping identifies 41 new loci associated with breast cancer risk. *Nature Genetics*, **2013**, 45, 353-61, 361e1-2	36.3	813
198	The International Neuroblastoma Pathology Classification (the Shimada system). *Cancer*, **1999**, 86, 364-372	6.4	781
197	Association analysis identifies 65 new breast cancer risk loci. *Nature*, **2017**, 551, 92-94	50.4	643
196	Multiple regions within 8q24 independently affect risk for prostate cancer. *Nature Genetics*, **2007**, 39, 638-44	36.3	563
195	Ethnic and racial differences in the smoking-related risk of lung cancer. *New England Journal of Medicine*, **2006**, 354, 333-42	59.2	538
194	Effects of inhalable particles on respiratory health of children. *The American Review of Respiratory Disease*, **1989**, 139, 587-94		509
193	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
192	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
191	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
190	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. *Nature Genetics*, **2014**, 46, 1103-9	36.3	331
189	Genome-wide association studies identify four ER negative-specific breast cancer risk loci. *Nature Genetics*, **2013**, 45, 392-8, 398e1-2	36.3	327
188	Exploiting gene-environment interaction to detect genetic associations. *Human Heredity*, **2007**, 63, 111-9	1.1	322
187	A common genetic risk factor for colorectal and prostate cancer. *Nature Genetics*, **2007**, 39, 954-6	36.3	304
186	Singapore Chinese Health Study: development, validation, and calibration of the quantitative food frequency questionnaire. *Nutrition and Cancer*, **2001**, 39, 187-95	2.8	299
185	Genetic analyses of diverse populations improves discovery for complex traits. *Nature*, **2019**, 570, 514-518	50.4	291
184	Heterogeneity of breast cancer associations with five susceptibility loci by clinical and pathological characteristics. *PLoS Genetics*, **2008**, 4, e1000054	6	280

183	A genome-wide association study identifies susceptibility loci for ovarian cancer at 2q31 and 8q24. *Nature Genetics*, **2010**, 42, 874-9	36.3	277
182	Metastatic sites in stage IV and IVS neuroblastoma correlate with age, tumor biology, and survival. *Journal of Pediatric Hematology/Oncology*, **1999**, 21, 181-9	1.2	274
181	A common variant at the TERT-CLPTM1L locus is associated with estrogen receptor-negative breast cancer. *Nature Genetics*, **2011**, 43, 1210-4	36.3	253
180	Meta-analysis: dietary fat intake, serum estrogen levels, and the risk of breast cancer. *Journal of the National Cancer Institute*, **1999**, 91, 529-34	9.7	242
179	A genome-wide association study identifies a new ovarian cancer susceptibility locus on 9p22.2. *Nature Genetics*, **2009**, 41, 996-1000	36.3	240
178	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
177	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
176	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
175	Common variants at 19p13 are associated with susceptibility to ovarian cancer. *Nature Genetics*, **2010**, 42, 880-4	36.3	210
174	Transferability of tag SNPs in genetic association studies in multiple populations. *Nature Genetics*, **2006**, 38, 1298-303	36.3	198
173	Favorable biology and outcome of stage IV-S neuroblastoma with supportive care or minimal therapy: a Children's Cancer Group study. *Journal of Clinical Oncology*, **2000**, 18, 477-86	2.2	195
172	Allelic deletion at 11q23 is common in MYCN single copy neuroblastomas. *Oncogene*, **1999**, 18, 4948-57	9.2	193
171	Revision of the International Neuroblastoma Pathology Classification: confirmation of favorable and unfavorable prognostic subsets in ganglioneuroblastoma, nodular. *Cancer*, **2003**, 98, 2274-81	6.4	189
170	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
169	Biologic variables in the outcome of stages I and II neuroblastoma treated with surgery as primary therapy: a children's cancer group study. *Journal of Clinical Oncology*, **2000**, 18, 18-26	2.2	183
168	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
167	Genome-wide association study of prostate cancer in men of African ancestry identifies a susceptibility locus at 17q21. *Nature Genetics*, **2011**, 43, 570-3	36.3	171
166	Functional variants at the 11q13 risk locus for breast cancer regulate cyclin D1 expression through long-range enhancers. *American Journal of Human Genetics*, **2013**, 92, 489-503	11	167

165	Biologic factors determine prognosis in infants with stage IV neuroblastoma: A prospective Children's Cancer Group study. *Journal of Clinical Oncology*, **2000**, 18, 1260-8	2.2	166
164	Tag SNP selection for association studies. *Genetic Epidemiology*, **2004**, 27, 365-74	2.6	149
163	Risk factors for renal cell cancer: the multiethnic cohort. *American Journal of Epidemiology*, **2007**, 166, 932-40	3.8	146
162	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, 5373-84	5.6	143
161	Genome-wide association study identifies new prostate cancer susceptibility loci. *Human Molecular Genetics*, **2011**, 20, 3867-75	5.6	143
160	Consistent association of type 2 diabetes risk variants found in europeans in diverse racial and ethnic groups. *PLoS Genetics*, **2010**, 6, e1001078	6	142
159	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, 1252-63	9.7	134
158	Genetic variation at the CYP19A1 locus predicts circulating estrogen levels but not breast cancer risk in postmenopausal women. *Cancer Research*, **2007**, 67, 1893-7	10.1	134
157	A comprehensive haplotype analysis of CYP19 and breast cancer risk: the Multiethnic Cohort. *Human Molecular Genetics*, **2003**, 12, 2679-92	5.6	131
156	Quantitative tumor cell content of bone marrow and blood as a predictor of outcome in stage IV neuroblastoma: a Children's Cancer Group Study. *Journal of Clinical Oncology*, **2000**, 18, 4067-76	2.2	130
155	Identification of subsets of neuroblastomas by combined histopathologic and N-myc analysis. *Journal of the National Cancer Institute*, **1995**, 87, 1470-6	9.7	128
154	Impact of radiotherapy for high-risk neuroblastoma: a Children's Cancer Group study. *International Journal of Radiation Oncology Biology Physics*, **2003**, 56, 28-39	4	125
153	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. *Nature Communications*, **2013**, 4, 1628	17.4	124
152	Loss of heterozygosity at 1p36 independently predicts for disease progression but not decreased overall survival probability in neuroblastoma patients: a Children's Cancer Group study. *Journal of Clinical Oncology*, **2000**, 18, 1888-99	2.2	124
151	Association of diabetes with prostate cancer risk in the multiethnic cohort. *American Journal of Epidemiology*, **2009**, 169, 937-45	3.8	110
150	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
149	Optimal two-stage genotyping designs for genome-wide association scans. *Genetic Epidemiology*, **2006**, 30, 356-68	2.6	107
148	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

147	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
146	Characterizing genetic risk at known prostate cancer susceptibility loci in African Americans. *PLoS Genetics*, **2011**, 7, e1001387	6	98
145	Common genetic variation in IGF1 and prostate cancer risk in the Multiethnic Cohort. *Journal of the National Cancer Institute*, **2006**, 98, 123-34	9.7	97
144	Generalizability of associations from prostate cancer genome-wide association studies in multiple populations. *Cancer Epidemiology Biomarkers and Prevention*, **2009**, 18, 1285-9	4	95
143	IGF-1, IGFBP-1, and IGFBP-3 polymorphisms predict circulating IGF levels but not breast cancer risk: findings from the Breast and Prostate Cancer Cohort Consortium (BPC3). *PLoS ONE*, **2008**, 3, e2578	3.7	93
142	Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. *Nature Communications*, **2014**, 4, 4999	17.4	87
141	Fine-scale mapping of the FGFR2 breast cancer risk locus: putative functional variants differentially bind FOXA1 and E2F1. *American Journal of Human Genetics*, **2013**, 93, 1046-60	11	80
140	Analysis of Repeated Ordered Categorical Outcomes with Possibly Missing Observations and Time-Dependent Covariates. *Journal of the American Statistical Association*, **1988**, 83, 631-637	2.8	78
139	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
138	Common genetic determinants of breast-cancer risk in East Asian women: a collaborative study of 23 637 breast cancer cases and 25 579 controls. *Human Molecular Genetics*, **2013**, 22, 2539-50	5.6	75
137	Dietary patterns and breast cancer risk in the California Teachers Study cohort. *American Journal of Clinical Nutrition*, **2013**, 98, 1524-32	7	75
136	TEMPORAL AGGREGATION IN THE ARIMA PROCESS. *Journal of Time Series Analysis*, **1986**, 7, 279-292	0.8	73
135	Prostate Cancer Susceptibility in Men of African Ancestry at 8q24. *Journal of the National Cancer Institute*, **2016**, 108,	9.7	72
134	Population distribution of lifetime risk of ovarian cancer in the United States. *Cancer Epidemiology Biomarkers and Prevention*, **2015**, 24, 671-676	4	67
133	Generalizability and epidemiologic characterization of eleven colorectal cancer GWAS hits in multiple populations. *Cancer Epidemiology Biomarkers and Prevention*, **2011**, 20, 70-81	4	66
132	A genome-wide association study of breast cancer in women of African ancestry. *Human Genetics*, **2013**, 132, 39-48	6.3	63
131	Trans-ethnic genome-wide association study of colorectal cancer identifies a new susceptibility locus in VTI1A. *Nature Communications*, **2014**, 5, 4613	17.4	62
130	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

129	Fine-mapping of breast cancer susceptibility loci characterizes genetic risk in African Americans. *Human Molecular Genetics*, **2011**, 20, 4491-503	5.6	58
128	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, 2468-75	10.1	58
127	Association of type 2 diabetes susceptibility variants with advanced prostate cancer risk in the Breast and Prostate Cancer Cohort Consortium. *American Journal of Epidemiology*, **2012**, 176, 1121-9	3.8	57
126	Eighteen insulin-like growth factor pathway genes, circulating levels of IGF-I and its binding protein, and risk of prostate and breast cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2010**, 19, 2877-87	4	54
125	Genetic variation in the HSD17B1 gene and risk of prostate cancer. *PLoS Genetics*, **2005**, 1, e68	6	54
124	Generalizability of established prostate cancer risk variants in men of African ancestry. *International Journal of Cancer*, **2015**, 136, 1210-7	7.5	51
123	Dietary assessment in the California Teachers Study: reproducibility and validity. *Cancer Causes and Control*, **2008**, 19, 595-603	2.8	51
122	Clarifying the PROGINS allele association in ovarian and breast cancer risk: a haplotype-based analysis. *Journal of the National Cancer Institute*, **2005**, 97, 51-9	9.7	51
121	Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. *Cancer Research*, **2017**, 77, 2789-2799	10.1	49
120	Prediction of breast cancer risk by genetic risk factors, overall and by hormone receptor status. *Journal of Medical Genetics*, **2012**, 49, 601-8	5.8	49
119	CYP17 genetic variation and risk of breast and prostate cancer from the National Cancer Institute Breast and Prostate Cancer Cohort Consortium (BPC3). *Cancer Epidemiology Biomarkers and Prevention*, **2007**, 16, 2237-46	4	49
118	Characterizing associations and SNP-environment interactions for GWAS-identified prostate cancer risk markers--results from BPC3. *PLoS ONE*, **2011**, 6, e17142	3.7	49
117	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
116	A haplotype-based case-control study of BRCA1 and sporadic breast cancer risk. *Cancer Research*, **2005**, 65, 7516-22	10.1	48
115	Common variation in BRCA2 and breast cancer risk: a haplotype-based analysis in the Multiethnic Cohort. *Human Molecular Genetics*, **2004**, 13, 2431-41	5.6	48
114	Multiple nonglycemic genomic loci are newly associated with blood level of glycated hemoglobin in East Asians. *Diabetes*, **2014**, 63, 2551-62	0.9	46
113	Evaluating genetic risk for prostate cancer among Japanese and Latinos. *Cancer Epidemiology Biomarkers and Prevention*, **2012**, 21, 2048-58	4	46
112	A comprehensive analysis of common IGF1, IGFBP1 and IGFBP3 genetic variation with prospective IGF-I and IGFBP-3 blood levels and prostate cancer risk among Caucasians. *Human Molecular Genetics*, **2010**, 19, 3089-101	5.6	46

111	The role of KRAS rs61764370 in invasive epithelial ovarian cancer: implications for clinical testing. *Clinical Cancer Research*, **2011**, 17, 3742-50	12.9	45
110	Propensity for Intra-abdominal and Hepatic Adiposity Varies Among Ethnic Groups. *Gastroenterology*, **2019**, 156, 966-975.e10	13.3	44
109	A controlled 2-mo dietary fat reduction and soy food supplementation study in postmenopausal women. *American Journal of Clinical Nutrition*, **2005**, 81, 1133-41	7	43
108	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
107	Evaluation of candidate stromal epithelial cross-talk genes identifies association between risk of serous ovarian cancer and TERT, a cancer susceptibility "hot-spot". *PLoS Genetics*, **2010**, 6, e1001016	6	42
106	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
105	Bayesian spatial modeling of haplotype associations. *Human Heredity*, **2003**, 56, 32-40	1.1	40
104	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. *Human Molecular Genetics*, **2010**, 19, 3873-84	5.6	39
103	Methodological Issues in Multistage Genome-wide Association Studies. *Statistical Science*, **2009**, 24, 414-429	2.4	39
102	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. *Journal of the National Cancer Institute*, **2017**, 109,	9.7	38
101	Comprehensive association testing of common genetic variation in DNA repair pathway genes in relationship with breast cancer risk in multiple populations. *Human Molecular Genetics*, **2008**, 17, 825-34	5.6	38
100	Haplotype-based association studies of IGFBP1 and IGFBP3 with prostate and breast cancer risk: the multiethnic cohort. *Cancer Epidemiology Biomarkers and Prevention*, **2006**, 15, 1993-7	4	38
99	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. *Nature Communications*, **2016**, 7, 10979	17.4	37
98	Combined and interactive effects of environmental and GWAS-identified risk factors in ovarian cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2013**, 22, 880-90	4	37
97	Recent breast cancer incidence trends according to hormone therapy use: the California Teachers Study cohort. *Breast Cancer Research*, **2010**, 12, R4	8.3	37
96	Association between invasive ovarian cancer susceptibility and 11 best candidate SNPs from breast cancer genome-wide association study. *Human Molecular Genetics*, **2009**, 18, 2297-304	5.6	37
95	Statistical Issues in Studies of the Long-Term Effects of Air Pollution: The Southern California Children's Health Study. *Statistical Science*, **2004**, 19, 414	2.4	37
94	Histopathology defines prognostic subsets of ganglioneuroblastoma, nodular. *Cancer*, **2000**, 89, 1150-1161	6.4	37

93	Genetic determinants of CYP2A6 activity across racial/ethnic groups with different risks of lung cancer and effect on their smoking intensity. *Carcinogenesis*, **2016**, 37, 269-279	4.6	36
92	Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. *Human Molecular Genetics*, **2015**, 24, 2966-84	5.6	36
91	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, 3749-57	5.6	36
90	Haplotypes of the estrogen receptor beta gene and breast cancer risk. *International Journal of Cancer*, **2008**, 122, 387-92	7.5	36
89	Racial/Ethnic Differences in Lung Cancer Incidence in the Multiethnic Cohort Study: An Update. *Journal of the National Cancer Institute*, **2019**, 111, 811-819	9.7	35
88	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
87	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
86	Pooled analysis of phosphatidylinositol 3-kinase pathway variants and risk of prostate cancer. *Cancer Research*, **2010**, 70, 2389-96	10.1	35
85	Deletion of 11q23 is a frequent event in the evolution of MYCN single-copy high-risk neuroblastomas. *Medical and Pediatric Oncology*, **2000**, 35, 544-6		35
84	Diabetes and racial/ethnic differences in hepatocellular carcinoma risk: the multiethnic cohort. *Journal of the National Cancer Institute*, **2014**, 106,	9.7	34
83	Levels of beta-microseminoprotein in blood and risk of prostate cancer in multiple populations. *Journal of the National Cancer Institute*, **2013**, 105, 237-43	9.7	34
82	A METHODOLOGICAL NOTE ON THE DISAGGREGATION OF TIME SERIES TOTALS. *Journal of Time Series Analysis*, **1986**, 7, 293-302	0.8	33
81	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
80	Self-reported ethnicity, genetic structure and the impact of population stratification in a multiethnic study. *Human Genetics*, **2010**, 128, 165-77	6.3	32
79	A comprehensive analysis of common genetic variation in prolactin (PRL) and PRL receptor (PRLR) genes in relation to plasma prolactin levels and breast cancer risk: the multiethnic cohort. *BMC Medical Genetics*, **2007**, 8, 72	2.1	32
78	European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. *Nature Communications*, **2020**, 11, 3833	17.4	31
77	Software for tag single nucleotide polymorphism selection. *Human Genomics*, **2005**, 2, 144-51	6.8	30
76	Enlarged and prominent nucleoli may be indicative of MYCN amplification: a study of neuroblastoma (Schwannian stroma-poor), undifferentiated/poorly differentiated subtype with high mitosis-karyorrhexis index. *Cancer*, **2005**, 103, 174-80	6.4	30

75	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, 1018-27	3.8	29
74	CYP19A1 genetic variation in relation to prostate cancer risk and circulating sex hormone concentrations in men from the Breast and Prostate Cancer Cohort Consortium. *Cancer Epidemiology Biomarkers and Prevention*, **2009**, 18, 2734-44	4	29
73	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	2.2	29
72	A comprehensive examination of breast cancer risk loci in African American women. *Human Molecular Genetics*, **2014**, 23, 5518-26	5.6	28
71	Sequence variants of estrogen receptor beta and risk of prostate cancer in the National Cancer Institute Breast and Prostate Cancer Cohort Consortium. *Cancer Epidemiology Biomarkers and Prevention*, **2007**, 16, 1973-81	4	28
70	A comprehensive analysis of the androgen receptor gene and risk of breast cancer: results from the National Cancer Institute Breast and Prostate Cancer Cohort Consortium (BPC3). *Breast Cancer Research*, **2006**, 8, R54	8.3	28
69	Association of CYP2A6 activity with lung cancer incidence in smokers: The multiethnic cohort study. *PLoS ONE*, **2017**, 12, e0178435	3.7	28
68	Tobacco biomarkers and genetic/epigenetic analysis to investigate ethnic/racial differences in lung cancer risk among smokers. *Npj Precision Oncology*, **2018**, 2, 17	9.8	25
67	Exploring genetic susceptibility to cancer in diverse populations. *Current Opinion in Genetics and Development*, **2010**, 20, 330-5	4.9	25
66	Common genetic variation at PTEN and risk of sporadic breast and prostate cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2006**, 15, 1021-5	4	25
65	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
64	Interethnic differences in pancreatic cancer incidence and risk factors: The Multiethnic Cohort. *Cancer Medicine*, **2019**, 8, 3592-3603	4.8	23
63	The role of local ancestry adjustment in association studies using admixed populations. *Genetic Epidemiology*, **2014**, 38, 502-15	2.6	23
62	Dietary patterns and risk of ovarian cancer in the California Teachers Study cohort. *Nutrition and Cancer*, **2008**, 60, 285-91	2.8	23
61	Association of Common Genetic Variants With Contralateral Breast Cancer Risk in the WECARE Study. *Journal of the National Cancer Institute*, **2017**, 109,	9.7	22
60	Association of internal smoking dose with blood DNA methylation in three racial/ethnic populations. *Clinical Epigenetics*, **2018**, 10, 110	7.7	22
59	Comparison of prostate-specific antigen and hormone levels among men in Singapore and the United States. *Cancer Epidemiology Biomarkers and Prevention*, **2005**, 14, 1692-6	4	22
58	Whole-exome sequencing of over 4100 men of African ancestry and prostate cancer risk. *Human Molecular Genetics*, **2016**, 25, 371-81	5.6	19

57	Igf-I genetic variation and breast cancer: the multiethnic cohort. *Cancer Epidemiology Biomarkers and Prevention*, **2006**, 15, 172-4	4	19
56	Genetic variation in insulin-like growth factor 2 may play a role in ovarian cancer risk. *Human Molecular Genetics*, **2011**, 20, 2263-72	5.6	18
55	Estimates of Radiation Effects on Cancer Risks in the Mayak Worker, Techa River and Atomic Bomb Survivor Studies. *Radiation Protection Dosimetry*, **2017**, 173, 26-31	0.9	17
54	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
53	A genome-wide scan for breast cancer risk haplotypes among African American women. *PLoS ONE*, **2013**, 8, e57298	3.7	17
52	Caution in generalizing known genetic risk markers for breast cancer across all ethnic/racial populations. *European Journal of Human Genetics*, **2011**, 19, 243-5	5.3	17
51	A systematic assessment of common genetic variation in CYP11A and risk of breast cancer. *Cancer Research*, **2006**, 66, 12019-25	10.1	17
50	Asymptotic equivalence between two score tests for haplotype-specific risk in general linear models. *Genetic Epidemiology*, **2005**, 29, 166-70	2.6	17
49	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
48	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
47	A common prostate cancer risk variant 5' of microseminoprotein-beta (MSMB) is a strong predictor of circulating beta-microseminoprotein (MSP) levels in multiple populations. *Cancer Epidemiology Biomarkers and Prevention*, **2010**, 19, 2639-46	4	16
46	Utilizing HapMap and tagging SNPs. *Methods in Molecular Medicine*, **2008**, 141, 37-54		16
45	Re: the use of inferred haplotypes in downstream analysis. *American Journal of Human Genetics*, **2007**, 81, 863-5; author reply 865-6	11	16
44	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
43	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
42	Multi-SNP Haplotype Analysis Methods for Association Analysis. *Methods in Molecular Biology*, **2017** , 1666, 485-504	1.4	15
41	Pleiotropy of cancer susceptibility variants on the risk of non-Hodgkin lymphoma: the PAGE consortium. *PLoS ONE*, **2014**, 9, e89791	3.7	14
40	Genetic variation in peroxisome proliferator-activated receptor gamma, soy, and mammographic density in Singapore Chinese women. *Cancer Epidemiology Biomarkers and Prevention*, **2012**, 21, 635-44	4	14

39	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
38	The PAGE Study: How Genetic Diversity Improves Our Understanding of the Architecture of Complex Traits		14
37	Characterizing Genetic Susceptibility to Breast Cancer in Women of African Ancestry. *Cancer Epidemiology Biomarkers and Prevention*, **2017**, 26, 1016-1026	4	12
36	Evaluating the ovarian cancer gonadotropin hypothesis: a candidate gene study. *Gynecologic Oncology*, **2015**, 136, 542-8	4.9	12
35	Associations Between Genetic Ancestries and Nicotine Metabolism Biomarkers in the Multiethnic Cohort Study. *American Journal of Epidemiology*, **2015**, 182, 945-51	3.8	11
34	Design, Analysis, and Interpretation of Genome-Wide Association Scans **2014**,		11
33	No association of type 2 diabetes risk variants and prostate cancer risk: the multiethnic cohort and PAGE. *Cancer Epidemiology Biomarkers and Prevention*, **2011**, 20, 1979-81	4	11
32	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
31	IGF2R missense single-nucleotide polymorphisms and breast cancer risk: the multiethnic cohort study. *Cancer Epidemiology Biomarkers and Prevention*, **2009**, 18, 1922-4	4	10
30	Disaggregation of Time Series Models. *Journal of the Royal Statistical Society Series B: Methodological*, **1990**, 52, 453-467		9
29	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
28	Genome-Wide Association Study of Liver Fat: The Multiethnic Cohort Adiposity Phenotype Study. *Hepatology Communications*, **2020**, 4, 1112-1123	6	8
27	Evaluation of 71 Coronary Artery Disease Risk Variants in a Multiethnic Cohort. *Frontiers in Cardiovascular Medicine*, **2018**, 5, 19	5.4	8
26	An eigenvalue approach to the limiting behavior of time series aggregates. *Annals of the Institute of Statistical Mathematics*, **1988**, 40, 101-110	1	8
25	Multi-SNP haplotype analysis methods for association analysis. *Methods in Molecular Biology*, **2012**, 850, 423-52	1.4	7
24	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
23	A splicing variant of TERT identified by GWAS interacts with menopausal estrogen therapy in risk of ovarian cancer. *International Journal of Cancer*, **2016**, 139, 2646-2654	7.5	6
22	An utter refutation of the "Fundamental Theorem of the HapMap" by Terwilliger and Hiekkalinna. *European Journal of Human Genetics*, **2006**, 14, 1238-9	5.3	6

21	Genetic variation in the HSD17B1 gene and risk of prostate cancer. *PLoS Genetics*, **2005**, preprint, e68	6	6
20	Meta-Analysis of Rare Variant Association Tests in Multiethnic Populations. *Genetic Epidemiology*, **2016**, 40, 57-65	2.6	6
19	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
18	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
17	Statistical issues in epidemiologic studies of the health effects of ambient acid aerosols. *Canadian Journal of Statistics*, **1988**, 16, 5-13	0.4	5
16	Genetic polymorphisms of the GNRH1 and GNRHR genes and risk of breast cancer in the National Cancer Institute Breast and Prostate Cancer Cohort Consortium (BPC3). *BMC Cancer*, **2009**, 9, 257	4.8	4
15	The Impact of GWAS Findings on Cancer Etiology and Prevention. *Current Epidemiology Reports*, **2014**, 1, 130-137	2.9	3
14	Breast Cancer Among Asian Americans **2016**, 187-218		3
13	RESPONSE: re: meta-analysis: dietary fat intake, serum estrogen levels, and the risk of breast cancer. *Journal of the National Cancer Institute*, **2000**, 92, 78A-78	9.7	2
12	Association between mitochondrial genetic variation and breast cancer risk: The Multiethnic Cohort. *PLoS ONE*, **2019**, 14, e0222284	3.7	1
11	Correcting for Hidden Population Structure in Single Marker Association Testing and Estimation **2014**, 135-181		1
10	Associations of the gut microbiome with hepatic adiposity in the Multiethnic Cohort Adiposity Phenotype Study. *Gut Microbes*, **2021**, 13, 1965463	8.8	1
9	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
8	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
7	Genome-wide association study of pancreatic fat: The Multiethnic Cohort Adiposity Phenotype Study. *PLoS ONE*, **2021**, 16, e0249615	3.7	0
6	Post-GWAS Analyses **2014**, 285-327		
5	Prostate cancer susceptibility polymorphism rs2660753 is not associated with invasive ovarian cancer. *Cancer Epidemiology Biomarkers and Prevention*, **2011**, 20, 1028-31	4	
4	Lung Cancer Among Asian Americans **2016**, 107-136		

List of Publications

3 SNP Imputation for Association Studies **2014**, 213-242

2 An Introduction to Association Analysis **2014**, 79-133

1 Haplotype Imputation for Association Analysis **2014**, 183-211