

# Daniel O Stram

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

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

| #   | Paper   | IF   | Citations |
|-----|---|------|-----------|
| 200 | 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> , <b>2022</b> ,                           | 11   | 1         |
| 199 | Associations of the gut microbiome with hepatic adiposity in the Multiethnic Cohort Adiposity Phenotype Study. <i>Gut Microbes</i> , <b>2021</b> , 13, 1965463  | 8.8  | 1         |
| 198 | Genome-wide association study of pancreatic fat: The Multiethnic Cohort Adiposity Phenotype Study. <i>PLoS ONE</i> , <b>2021</b> , 16, e0249615   | 3.7  | 0         |
| 197 | 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> , <b>2020</b> , 16, e1008684                               | 6    | 5         |
| 196 | Genome-Wide Association Study of Liver Fat: The Multiethnic Cohort Adiposity Phenotype Study. <i>Hepatology Communications</i> , <b>2020</b> , 4, 1112-1123   | 6    | 8         |
| 195 | Urinary N7-(1-hydroxy-3-buten-2-yl) guanine adducts in humans: temporal stability and association with smoking. <i>Mutagenesis</i> , <b>2020</b> , 35, 19-26  | 2.8  | 9         |
| 194 | European polygenic risk score for prediction of breast cancer shows similar performance in Asian women. <i>Nature Communications</i> , <b>2020</b> , 11, 3833   | 17.4 | 31        |
| 193 | Association between mitochondrial genetic variation and breast cancer risk: The Multiethnic Cohort. <i>PLoS ONE</i> , <b>2019</b> , 14, e0222284  | 3.7  | 1         |
| 192 | Racial/Ethnic Differences in Lung Cancer Incidence in the Multiethnic Cohort Study: An Update. <i>Journal of the National Cancer Institute</i> , <b>2019</b> , 111, 811-819   | 9.7  | 35        |
| 191 | Genetic analyses of diverse populations improves discovery for complex traits. <i>Nature</i> , <b>2019</b> , 570, 514-518   | 50.4 | 291       |
| 190 | Interethnic differences in pancreatic cancer incidence and risk factors: The Multiethnic Cohort. <i>Cancer Medicine</i> , <b>2019</b> , 8, 3592-3603  | 4.8  | 23        |
| 189 | Propensity for Intra-abdominal and Hepatic Adiposity Varies Among Ethnic Groups. <i>Gastroenterology</i> , <b>2019</b> , 156, 966-975.e10   | 13.3 | 44        |
| 188 | Evaluation of 71 Coronary Artery Disease Risk Variants in a Multiethnic Cohort. <i>Frontiers in Cardiovascular Medicine</i> , <b>2018</b> , 5, 19   | 5.4  | 8         |
| 187 | Association of internal smoking dose with blood DNA methylation in three racial/ethnic populations. <i>Clinical Epigenetics</i> , <b>2018</b> , 10, 110   | 7.7  | 22        |
| 186 | Tobacco biomarkers and genetic/epigenetic analysis to investigate ethnic/racial differences in lung cancer risk among smokers. <i>Npj Precision Oncology</i> , <b>2018</b> , 2, 17  | 9.8  | 25        |
| 185 | Breast Cancer Family History and Contralateral Breast Cancer Risk in Young Women: An Update From the Women's Environmental Cancer and Radiation Epidemiology Study. <i>Journal of Clinical Oncology</i> , <b>2018</b> , 36, 1513-1520 | 2.2  | 29        |
| 184 | Growth factor genes and change in mammographic density after stopping combined hormone therapy in the California Teachers Study. <i>BMC Cancer</i> , <b>2018</b> , 18, 1072   | 4.8  | 1         |

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| 183 | Estimates of Radiation Effects on Cancer Risks in the Mayak Worker, Techa River and Atomic Bomb Survivor Studies. <i>Radiation Protection Dosimetry</i> , <b>2017</b> , 173, 26-31  | 0.9  | 17  |
| 182 | Coffee Drinking and Alcoholic and Nonalcoholic Fatty Liver Diseases and Viral Hepatitis in the Multiethnic Cohort. <i>Clinical Gastroenterology and Hepatology</i> , <b>2017</b> , 15, 1305-1307                              | 6.9  | 17  |
| 181 | Hypomorphic Missense Variants Confer Moderate Risks of Breast Cancer. <i>Cancer Research</i> , <b>2017</b> , 77, 2789-2799  | 10.1 | 49  |
| 180 | Association of Common Genetic Variants With Contralateral Breast Cancer Risk in the WECARE Study. <i>Journal of the National Cancer Institute</i> , <b>2017</b> , 109,  | 9.7  | 22  |
| 179 | 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> , <b>2017</b> , 26, 1034-1042   | 4    | 15  |
| 178 | Characterizing Genetic Susceptibility to Breast Cancer in Women of African Ancestry. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2017</b> , 26, 1016-1026   | 4    | 12  |
| 177 | Multi-SNP Haplotype Analysis Methods for Association Analysis. <i>Methods in Molecular Biology</i> , <b>2017</b> , 1666, 485-504  | 1.4  | 15  |
| 176 | Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , <b>2017</b> , 551, 92-94  | 50.4 | 643 |
| 175 | Impact of common genetic determinants of Hemoglobin A1c on type 2 diabetes risk and diagnosis in ancestrally diverse populations: A transeethnic genome-wide meta-analysis. <i>PLoS Medicine</i> , <b>2017</b> , 14, e1002383 | 11.6 | 223 |
| 174 | Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. <i>Journal of the National Cancer Institute</i> , <b>2017</b> , 109,  | 9.7  | 38  |
| 173 | Association of CYP2A6 activity with lung cancer incidence in smokers: The multiethnic cohort study. <i>PLoS ONE</i> , <b>2017</b> , 12, e0178435  | 3.7  | 28  |
| 172 | A splicing variant of TERT identified by GWAS interacts with menopausal estrogen therapy in risk of ovarian cancer. <i>International Journal of Cancer</i> , <b>2016</b> , 139, 2646-2654                                     | 7.5  | 6   |
| 171 | Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. <i>Cancer Discovery</i> , <b>2016</b> , 6, 1052-67       | 24.4 | 104 |
| 170 | Fine scale mapping of the 17q22 breast cancer locus using dense SNPs, genotyped within the Collaborative Oncological Gene-Environment Study (COGs). <i>Scientific Reports</i> , <b>2016</b> , 6, 32512                        | 4.9  | 16  |
| 169 | Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. <i>Nature Communications</i> , <b>2016</b> , 7, 10979  | 17.4 | 37  |
| 168 | Prostate Cancer Susceptibility in Men of African Ancestry at 8q24. <i>Journal of the National Cancer Institute</i> , <b>2016</b> , 108,   | 9.7  | 72  |
| 167 | Genetic determinants of CYP2A6 activity across racial/ethnic groups with different risks of lung cancer and effect on their smoking intensity. <i>Carcinogenesis</i> , <b>2016</b> , 37, 269-279                              | 4.6  | 36  |
| 166 | Whole-exome sequencing of over 4100 men of African ancestry and prostate cancer risk. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 371-81  | 5.6  | 19  |

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| 165 | Benzene Uptake and Glutathione S-transferase T1 Status as Determinants of S-Phenylmercapturic Acid in Cigarette Smokers in the Multiethnic Cohort. <i>PLoS ONE</i> , <b>2016</b> , 11, e0150641   | 3.7  | 16  |
| 164 | Breast Cancer Among Asian Americans <b>2016</b> , 187-218   |      | 3   |
| 163 | Lung Cancer Among Asian Americans <b>2016</b> , 107-136   |      |     |
| 162 | Metabolites of the Polycyclic Aromatic Hydrocarbon Phenanthrene in the Urine of Cigarette Smokers from Five Ethnic Groups with Differing Risks for Lung Cancer. <i>PLoS ONE</i> , <b>2016</b> , 11, e0156203  | 3.7  | 17  |
| 161 | Prevalence of chronic liver disease and cirrhosis by underlying cause in understudied ethnic groups: The multiethnic cohort. <i>Hepatology</i> , <b>2016</b> , 64, 1969-1977  | 11.2 | 175 |
| 160 | Breast Cancer Risk From Modifiable and Nonmodifiable Risk Factors Among White Women in the United States. <i>JAMA Oncology</i> , <b>2016</b> , 2, 1295-1302   | 13.4 | 189 |
| 159 | Meta-Analysis of Rare Variant Association Tests in Multiethnic Populations. <i>Genetic Epidemiology</i> , <b>2016</b> , 40, 57-65   | 2.6  | 6   |
| 158 | Genome-wide association analysis of more than 120,000 individuals identifies 15 new susceptibility loci for breast cancer. <i>Nature Genetics</i> , <b>2015</b> , 47, 373-80  | 36.3 | 406 |
| 157 | Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. <i>Clinical Cancer Research</i> , <b>2015</b> , 21, 5264-76  | 12.9 | 24  |
| 156 | Integration of multiethnic fine-mapping and genomic annotation to prioritize candidate functional SNPs at prostate cancer susceptibility regions. <i>Human Molecular Genetics</i> , <b>2015</b> , 24, 5603-18                                       | 5.6  | 35  |
| 155 | Evaluating the ovarian cancer gonadotropin hypothesis: a candidate gene study. <i>Gynecologic Oncology</i> , <b>2015</b> , 136, 542-8   | 4.9  | 12  |
| 154 | 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. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 561-9 | 4    | 32  |
| 153 | Associations Between Genetic Ancestries and Nicotine Metabolism Biomarkers in the Multiethnic Cohort Study. <i>American Journal of Epidemiology</i> , <b>2015</b> , 182, 945-51   | 3.8  | 11  |
| 152 | Population distribution of lifetime risk of ovarian cancer in the United States. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 671-676   | 4    | 67  |
| 151 | Identification and characterization of novel associations in the CASP8/ALS2CR12 region on chromosome 2 with breast cancer risk. <i>Human Molecular Genetics</i> , <b>2015</b> , 24, 285-98  | 5.6  | 35  |
| 150 | ABO blood group alleles and prostate cancer risk: Results from the breast and prostate cancer cohort consortium (BPC3). <i>Prostate</i> , <b>2015</b> , 75, 1677-81   | 4.2  | 10  |
| 149 | Fine-mapping identifies two additional breast cancer susceptibility loci at 9q31.2. <i>Human Molecular Genetics</i> , <b>2015</b> , 24, 2966-84   | 5.6  | 36  |
| 148 | The contribution of common genetic variation to nicotine and cotinine glucuronidation in multiple ethnic/racial populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 119-27  | 4    | 42  |

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| 147 | Generalizability of established prostate cancer risk variants in men of African ancestry. <i>International Journal of Cancer</i> , <b>2015</b> , 136, 1210-7  | 7.5  | 51  |
| 146 | Fine-scale mapping of the 5q11.2 breast cancer locus reveals at least three independent risk variants regulating MAP3K1. <i>American Journal of Human Genetics</i> , <b>2015</b> , 96, 5-20   | 11   | 59  |
| 145 | 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> , <b>2015</b> , 10, e0124841   | 3.7  | 42  |
| 144 | Design, Analysis, and Interpretation of Genome-Wide Association Scans <b>2014</b> ,   |      | 11  |
| 143 | The Impact of GWAS Findings on Cancer Etiology and Prevention. <i>Current Epidemiology Reports</i> , <b>2014</b> , 1, 130-137   | 2.9  | 3   |
| 142 | A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. <i>Nature Genetics</i> , <b>2014</b> , 46, 1103-9  | 36.3 | 331 |
| 141 | Evidence that breast cancer risk at the 2q35 locus is mediated through IGFBP5 regulation. <i>Nature Communications</i> , <b>2014</b> , 4, 4999  | 17.4 | 87  |
| 140 | Fine-mapping IGF1 and prostate cancer risk in African Americans: the multiethnic cohort study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2014</b> , 23, 1928-32   | 4    | 5   |
| 139 | Pleiotropy of cancer susceptibility variants on the risk of non-Hodgkin lymphoma: the PAGE consortium. <i>PLoS ONE</i> , <b>2014</b> , 9, e89791  | 3.7  | 14  |
| 138 | Multiple nonglycemic genomic loci are newly associated with blood level of glycated hemoglobin in East Asians. <i>Diabetes</i> , <b>2014</b> , 63, 2551-62  | 0.9  | 46  |
| 137 | Nicotine N-glucuronidation relative to N-oxidation and C-oxidation and UGT2B10 genotype in five ethnic/racial groups. <i>Carcinogenesis</i> , <b>2014</b> , 35, 2526-33   | 4.6  | 103 |
| 136 | Additive interactions between susceptibility single-nucleotide polymorphisms identified in genome-wide association studies and breast cancer risk factors in the Breast and Prostate Cancer Cohort Consortium. <i>American Journal of Epidemiology</i> , <b>2014</b> , 180, 1018-27 | 3.8  | 29  |
| 135 | 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> , <b>2014</b> , 23, 6616-33  | 5.6  | 77  |
| 134 | Common non-synonymous SNPs associated with breast cancer susceptibility: findings from the Breast Cancer Association Consortium. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 6096-111   | 5.6  | 48  |
| 133 | Trans-ethnic genome-wide association study of colorectal cancer identifies a new susceptibility locus in VT11A. <i>Nature Communications</i> , <b>2014</b> , 5, 4613  | 17.4 | 62  |
| 132 | The role of local ancestry adjustment in association studies using admixed populations. <i>Genetic Epidemiology</i> , <b>2014</b> , 38, 502-15  | 2.6  | 23  |
| 131 | Post-GWAS Analyses <b>2014</b> , 285-327  |      |     |
| 130 | A comprehensive examination of breast cancer risk loci in African American women. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 5518-26   | 5.6  | 28  |

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| 129 | Diabetes and racial/ethnic differences in hepatocellular carcinoma risk: the multiethnic cohort. <i>Journal of the National Cancer Institute</i> , <b>2014</b> , 106,                                       | 9.7  | 34  |
| 128 | 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> , <b>2014</b> , 9, e87762             | 3.7  | 14  |
| 127 | SNP Imputation for Association Studies <b>2014</b> , 213-242  |      |     |
| 126 | Correcting for Hidden Population Structure in Single Marker Association Testing and Estimation <b>2014</b> , 135-181  |      | 1   |
| 125 | An Introduction to Association Analysis <b>2014</b> , 79-133  |      |     |
| 124 | Haplotype Imputation for Association Analysis <b>2014</b> , 183-211   |      |     |
| 123 | Fine-scale mapping of the FGFR2 breast cancer risk locus: putative functional variants differentially bind FOXA1 and E2F1. <i>American Journal of Human Genetics</i> , <b>2013</b> , 93, 1046-60            | 11   | 80  |
| 122 | Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. <i>Nature Genetics</i> , <b>2013</b> , 45, 371-84, 384e1-2                      | 36.3 | 422 |
| 121 | A genome-wide association study of breast cancer in women of African ancestry. <i>Human Genetics</i> , <b>2013</b> , 132, 39-48   | 6.3  | 63  |
| 120 | Functional variants at the 11q13 risk locus for breast cancer regulate cyclin D1 expression through long-range enhancers. <i>American Journal of Human Genetics</i> , <b>2013</b> , 92, 489-503             | 11   | 167 |
| 119 | Genome-wide association studies identify four ER negative-specific breast cancer risk loci. <i>Nature Genetics</i> , <b>2013</b> , 45, 392-8, 398e1-2   | 36.3 | 327 |
| 118 | Large-scale genotyping identifies 41 new loci associated with breast cancer risk. <i>Nature Genetics</i> , <b>2013</b> , 45, 353-61, 361e1-2  | 36.3 | 813 |
| 117 | Common genetic determinants of breast-cancer risk in East Asian women: a collaborative study of 23 637 breast cancer cases and 25 579 controls. <i>Human Molecular Genetics</i> , <b>2013</b> , 22, 2539-50 | 5.6  | 75  |
| 116 | Levels of beta-microseminoprotein in blood and risk of prostate cancer in multiple populations. <i>Journal of the National Cancer Institute</i> , <b>2013</b> , 105, 237-43                                 | 9.7  | 34  |
| 115 | Combined and interactive effects of environmental and GWAS-identified risk factors in ovarian cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2013</b> , 22, 880-90                       | 4    | 37  |
| 114 | Dietary patterns and breast cancer risk in the California Teachers Study cohort. <i>American Journal of Clinical Nutrition</i> , <b>2013</b> , 98, 1524-32  | 7    | 75  |
| 113 | Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. <i>Nature Communications</i> , <b>2013</b> , 4, 1628                                     | 17.4 | 124 |
| 112 | A genome-wide scan for breast cancer risk haplotypes among African American women. <i>PLoS ONE</i> , <b>2013</b> , 8, e57298  | 3.7  | 17  |

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| 111 | Evaluating genetic risk for prostate cancer among Japanese and Latinos. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2012</b> , 21, 2048-58   | 4    | 46  |
| 110 | Multi-SNP haplotype analysis methods for association analysis. <i>Methods in Molecular Biology</i> , <b>2012</b> , 850, 423-52   | 1.4  | 7   |
| 109 | A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. <i>Human Molecular Genetics</i> , <b>2012</b> , 21, 5373-84                        | 5.6  | 143 |
| 108 | Association of type 2 diabetes susceptibility variants with advanced prostate cancer risk in the Breast and Prostate Cancer Cohort Consortium. <i>American Journal of Epidemiology</i> , <b>2012</b> , 176, 1121-9 | 3.8  | 57  |
| 107 | Prediction of breast cancer risk by genetic risk factors, overall and by hormone receptor status. <i>Journal of Medical Genetics</i> , <b>2012</b> , 49, 601-8   | 5.8  | 49  |
| 106 | Genetic variation in peroxisome proliferator-activated receptor gamma, soy, and mammographic density in Singapore Chinese women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2012</b> , 21, 635-44   | 4    | 14  |
| 105 | Generalizability and epidemiologic characterization of eleven colorectal cancer GWAS hits in multiple populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2011</b> , 20, 70-81                  | 4    | 66  |
| 104 | Genome-wide association study of prostate cancer in men of African ancestry identifies a susceptibility locus at 17q21. <i>Nature Genetics</i> , <b>2011</b> , 43, 570-3   | 36.3 | 171 |
| 103 | Caution in generalizing known genetic risk markers for breast cancer across all ethnic/racial populations. <i>European Journal of Human Genetics</i> , <b>2011</b> , 19, 243-5                                     | 5.3  | 17  |
| 102 | Interactions between genetic variants and breast cancer risk factors in the breast and prostate cancer cohort consortium. <i>Journal of the National Cancer Institute</i> , <b>2011</b> , 103, 1252-63             | 9.7  | 134 |
| 101 | A common variant at the TERT-CLPTM1L locus is associated with estrogen receptor-negative breast cancer. <i>Nature Genetics</i> , <b>2011</b> , 43, 1210-4  | 36.3 | 253 |
| 100 | Genetic variation in insulin-like growth factor 2 may play a role in ovarian cancer risk. <i>Human Molecular Genetics</i> , <b>2011</b> , 20, 2263-72  | 5.6  | 18  |
| 99  | Genome-wide association study identifies new prostate cancer susceptibility loci. <i>Human Molecular Genetics</i> , <b>2011</b> , 20, 3867-75  | 5.6  | 143 |
| 98  | No association of type 2 diabetes risk variants and prostate cancer risk: the multiethnic cohort and PAGE. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2011</b> , 20, 1979-81                        | 4    | 11  |
| 97  | Prostate cancer susceptibility polymorphism rs2660753 is not associated with invasive ovarian cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2011</b> , 20, 1028-31                             | 4    |     |
| 96  | The role of KRAS rs61764370 in invasive epithelial ovarian cancer: implications for clinical testing. <i>Clinical Cancer Research</i> , <b>2011</b> , 17, 3742-50  | 12.9 | 45  |
| 95  | Fine-mapping of breast cancer susceptibility loci characterizes genetic risk in African Americans. <i>Human Molecular Genetics</i> , <b>2011</b> , 20, 4491-503  | 5.6  | 58  |
| 94  | Characterizing genetic risk at known prostate cancer susceptibility loci in African Americans. <i>PLoS Genetics</i> , <b>2011</b> , 7, e1001387  | 6    | 98  |



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| 93 | Characterizing associations and SNP-environment interactions for GWAS-identified prostate cancer risk markers--results from BPC3. <i>PLoS ONE</i> , <b>2011</b> , 6, e17142   | 3.7  | 49  |
| 92 | Common variants at 19p13 are associated with susceptibility to ovarian cancer. <i>Nature Genetics</i> , <b>2010</b> , 42, 880-4   | 36.3 | 210 |
| 91 | A genome-wide association study identifies susceptibility loci for ovarian cancer at 2q31 and 8q24. <i>Nature Genetics</i> , <b>2010</b> , 42, 874-9  | 36.3 | 277 |
| 90 | Eighteen insulin-like growth factor pathway genes, circulating levels of IGF-I and its binding protein, and risk of prostate and breast cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2010</b> , 19, 2877-84  | 4    | 54  |
| 89 | Pooled analysis of phosphatidylinositol 3-kinase pathway variants and risk of prostate cancer. <i>Cancer Research</i> , <b>2010</b> , 70, 2389-96   | 10.1 | 35  |
| 88 | Comprehensive analysis of common genetic variation in 61 genes related to steroid hormone and insulin-like growth factor-I metabolism and breast cancer risk in the NCI breast and prostate cancer cohort consortium. <i>Human Molecular Genetics</i> , <b>2010</b> , 19, 3873-84 | 5.6  | 39  |
| 87 | A common prostate cancer risk variant 5' of microseminoprotein-beta (MSMB) is a strong predictor of circulating beta-microseminoprotein (MSP) levels in multiple populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2010</b> , 19, 2639-46                    | 4    | 16  |
| 86 | 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. <i>Human Molecular Genetics</i> , <b>2010</b> , 19, 3089-101  | 5.6  | 46  |
| 85 | Evaluation of candidate stromal epithelial cross-talk genes identifies association between risk of serous ovarian cancer and TERT, a cancer susceptibility "hot-spot". <i>PLoS Genetics</i> , <b>2010</b> , 6, e1001016   | 6    | 42  |
| 84 | Consistent association of type 2 diabetes risk variants found in europeans in diverse racial and ethnic groups. <i>PLoS Genetics</i> , <b>2010</b> , 6, e1001078  | 6    | 142 |
| 83 | Exploring genetic susceptibility to cancer in diverse populations. <i>Current Opinion in Genetics and Development</i> , <b>2010</b> , 20, 330-5   | 4.9  | 25  |
| 82 | Recent breast cancer incidence trends according to hormone therapy use: the California Teachers Study cohort. <i>Breast Cancer Research</i> , <b>2010</b> , 12, R4  | 8.3  | 37  |
| 81 | Self-reported ethnicity, genetic structure and the impact of population stratification in a multiethnic study. <i>Human Genetics</i> , <b>2010</b> , 128, 165-77  | 6.3  | 32  |
| 80 | Methodological Issues in Multistage Genome-wide Association Studies. <i>Statistical Science</i> , <b>2009</b> , 24, 414-429   | 4.4  | 39  |
| 79 | Generalizability of associations from prostate cancer genome-wide association studies in multiple populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2009</b> , 18, 1285-9  | 4    | 95  |
| 78 | Association between invasive ovarian cancer susceptibility and 11 best candidate SNPs from breast cancer genome-wide association study. <i>Human Molecular Genetics</i> , <b>2009</b> , 18, 2297-304  | 5.6  | 37  |
| 77 | Association of diabetes with prostate cancer risk in the multiethnic cohort. <i>American Journal of Epidemiology</i> , <b>2009</b> , 169, 937-45  | 3.8  | 110 |
| 76 | IGF2R missense single-nucleotide polymorphisms and breast cancer risk: the multiethnic cohort study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2009</b> , 18, 1922-4  | 4    | 10  |



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| 75 | Quantitative trait loci predicting circulating sex steroid hormones in men from the NCI-Breast and Prostate Cancer Cohort Consortium (BPC3). <i>Human Molecular Genetics</i> , <b>2009</b> , 18, 3749-57  | 5.6  | 36  |
| 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. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2009</b> , 18, 2734-44 | 4    | 29  |
| 73 | 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). <i>BMC Cancer</i> , <b>2009</b> , 9, 257   | 4.8  | 4   |
| 72 | A genome-wide association study identifies a new ovarian cancer susceptibility locus on 9p22.2. <i>Nature Genetics</i> , <b>2009</b> , 41, 996-1000   | 36.3 | 240 |
| 71 | Utilizing HapMap and tagging SNPs. <i>Methods in Molecular Medicine</i> , <b>2008</b> , 141, 37-54  |      | 16  |
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