

# David V Conti

## List of Publications by Year in Descending Order

**Source:** <https://exaly.com/author-pdf/1883155/david-v-conti-publications-by-year.pdf>

**Version:** 2024-04-20

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

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

143  
papers

5,063  
citations

37  
h-index

68  
g-index

171  
ext. papers

6,798  
ext. citations

7.2  
avg, IF

4.68  
L-index

#	Paper	IF	Citations
143	Ancestral diversity improves discovery and fine-mapping of genetic loci for anthropometric traits-The Hispanic/Latino Anthropometry Consortium.. <i>Human Genetics and Genomics Advances</i> , <b>2022</b> , 3, 100099	0.8	0
142	Beyond GWAS of Colorectal Cancer: Evidence of Interaction with Alcohol Consumption and Putative Causal Variant for the 10q24.2 Region.. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2022</b> , OF1-OF13	4	0
141	Plasma concentrations of lipophilic persistent organic pollutants and glucose homeostasis in youth populations.. <i>Environmental Research</i> , <b>2022</b> , 212, 113296	7.9	1
140	Exposure to per- and Polyfluoroalkyl Substances and Markers of Liver Injury: A Systematic Review and Meta-Analysis.. <i>Environmental Health Perspectives</i> , <b>2022</b> , 130, 46001	8.4	8
139	Characteristics associated with COVID-19 vaccination status among staff and faculty of a large, diverse University in Los Angeles: The Trojan Pandemic Response Initiative.. <i>Preventive Medicine Reports</i> , <b>2022</b> , 27, 101802	2.6	1
138	Whole-Exome Sequencing in Multiplex Families to Identify Novel AYA Classical Hodgkin Lymphoma Predisposition Genes. <i>Blood</i> , <b>2021</b> , 138, 3499-3499	2.2	0
137	Differential Gene Expression in Circulating T-Cells in Long-Term Adolescent/Young Adult Hodgkin Lymphoma (AYAHL) Survivors and Their Unaffected Twins. <i>Blood</i> , <b>2021</b> , 138, 1332-1332	2.2	
136	Prenatal exposure to persistent organic pollutants and childhood obesity: A systematic review and meta-analysis of human studies. <i>Obesity Reviews</i> , <b>2021</b> , e13383	10.6	8
135	Advancing tools for human early lifecourse exposome research and translation (ATHLETE): Project overview.. <i>Environmental Epidemiology</i> , <b>2021</b> , 5, e166	0.2	2
134	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. <i>American Journal of Human Genetics</i> , <b>2021</b> , 108, 564-582	11	7
133	Evaluation of a Multiethnic Polygenic Risk Score Model for Prostate Cancer. <i>Journal of the National Cancer Institute</i> , <b>2021</b> ,	9.7	3
132	Prenatal and childhood exposure to air pollution and traffic and the risk of liver injury in European children. <i>Environmental Epidemiology</i> , <b>2021</b> , 5, e153	0.2	0
131	Bayesian variable selection with a pleiotropic loss function in Mendelian randomization. <i>Statistics in Medicine</i> , <b>2021</b> , 40, 5025-5045	2.3	1
130	An integrated risk and epidemiological model to estimate risk-stratified COVID-19 outcomes for Los Angeles County: March 1, 2020-March 1, 2021. <i>PLoS ONE</i> , <b>2021</b> , 16, e0253549	3.7	1
129	Association of Genetic Risk Score With NAFLD in An Ethnically Diverse Cohort. <i>Hepatology Communications</i> , <b>2021</b> , 5, 1689-1703	6	0
128	Cross-ancestry GWAS meta-analysis identifies six breast cancer loci in African and European ancestry women. <i>Nature Communications</i> , <b>2021</b> , 12, 4198	17.4	1
127	Identifying Novel Susceptibility Genes for Colorectal Cancer Risk From a Transcriptome-Wide Association Study of 125,478 Subjects. <i>Gastroenterology</i> , <b>2021</b> , 160, 1164-1178.e6	13.3	15

126	Germline Sequencing DNA Repair Genes in 5545 Men With Aggressive and Nonaggressive Prostate Cancer. <i>Journal of the National Cancer Institute</i> , <b>2021</b> , 113, 616-625	9.7	14
125	Prenatal metal mixtures and child blood pressure in the Rhea mother-child cohort in Greece. <i>Environmental Health</i> , <b>2021</b> , 20, 1	6	14
124	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. <i>Nature Genetics</i> , <b>2021</b> , 53, 65-75	36.3	62
123	Sign-based Shrinkage Based on an Asymmetric LASSO Penalty.. <i>Journal of Data Science</i> , <b>2021</b> , 19, 429-449	1	1
122	A Hierarchical Approach Using Marginal Summary Statistics for Multiple Intermediates in a Mendelian Randomization or Transcriptome Analysis. <i>American Journal of Epidemiology</i> , <b>2021</b> , 190, 1148-1158	3.8	1
121	Genetic architectures of proximal and distal colorectal cancer are partly distinct. <i>Gut</i> , <b>2021</b> , 70, 1325-1334	4.2	7
120	Rare Variants in the DNA Repair Pathway and the Risk of Colorectal Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2021</b> , 30, 895-903	4	1
119	In Utero Exposure to Mercury Is Associated With Increased Susceptibility to Liver Injury and Inflammation in Childhood. <i>Hepatology</i> , <b>2021</b> , 74, 1546-1559	11.2	3
118	Combined Effect of a Polygenic Risk Score and Rare Genetic Variants on Prostate Cancer Risk. <i>European Urology</i> , <b>2021</b> , 80, 134-138	10.2	7
117	Exposure to Perfluoroalkyl Substances and Glucose Homeostasis in Youth. <i>Environmental Health Perspectives</i> , <b>2021</b> , 129, 97002	8.4	3
116	Novel strategy for disease risk prediction incorporating predicted gene expression and DNA methylation data: a multi-phased study of prostate cancer. <i>Cancer Communications</i> , <b>2021</b> ,	9.4	1
115	Prenatal and postnatal exposure to PFAS and cardiometabolic factors and inflammation status in children from six European cohorts. <i>Environment International</i> , <b>2021</b> , 157, 106853	12.9	4
114	The Four-Kallikrein Panel Is Effective in Identifying Aggressive Prostate Cancer in a Multiethnic Population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2020</b> , 29, 1381-1388	4	15
113	A Germline Variant at 8q24 Contributes to Familial Clustering of Prostate Cancer in Men of African Ancestry. <i>European Urology</i> , <b>2020</b> , 78, 316-320	10.2	13
112	Variability in Cytogenetic Testing for Multiple Myeloma: A Comprehensive Analysis From Across the United States. <i>JCO Oncology Practice</i> , <b>2020</b> , 16, e1169-e1180	2.3	2
111	Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2020</b> , 29, 860-870	4	12
110	Functional informed genome-wide interaction analysis of body mass index, diabetes and colorectal cancer risk. <i>Cancer Medicine</i> , <b>2020</b> , 9, 3563-3573	4.8	4
109	Infectious mononucleosis, immune genotypes, and non-Hodgkin lymphoma (NHL): an InterLymph Consortium study. <i>Cancer Causes and Control</i> , <b>2020</b> , 31, 451-462	2.8	2

108	Pathogenic Variants in Cancer Predisposition Genes and Prostate Cancer Risk in Men of African Ancestry. <i>JCO Precision Oncology</i> , <b>2020</b> , 4, 32-43	3.6	13
107	A meta-analysis of genome-wide association studies of multiple myeloma among men and women of African ancestry. <i>Blood Advances</i> , <b>2020</b> , 4, 181-190	7.8	5
106	Cumulative Burden of Colorectal Cancer-Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. <i>Gastroenterology</i> , <b>2020</b> , 158, 1274-1286.e12	13.3	47
105	Interaction effect of alcohol consumption and Alzheimer disease polygenic risk score on the brain cortical thickness of cognitively normal subjects. <i>Alcohol</i> , <b>2020</b> , 85, 1-12	2.7	9
104	Perfluoroalkyl substances and severity of nonalcoholic fatty liver in Children: An untargeted metabolomics approach. <i>Environment International</i> , <b>2020</b> , 134, 105220	12.9	48
103	Replication and Genetic Risk Score Analysis for Pancreatic Cancer in a Diverse Multiethnic Population. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2020</b> , 29, 2686-2692	4	3
102	Environmental chemical burden in metabolic tissues and systemic biological pathways in adolescent bariatric surgery patients: A pilot untargeted metabolomic approach. <i>Environment International</i> , <b>2020</b> , 143, 105957	12.9	11
101	A Cross-Sectional Study Examining the Seroprevalence of Severe Acute Respiratory Syndrome Coronavirus 2 Antibodies in a University Student Population. <i>Journal of Adolescent Health</i> , <b>2020</b> , 67, 763-768	5.8	15
100	An integrative multi-omics analysis to identify candidate DNA methylation biomarkers related to prostate cancer risk. <i>Nature Communications</i> , <b>2020</b> , 11, 3905	17.4	12
99	Prenatal Exposure to Perfluoroalkyl Substances Associated With Increased Susceptibility to Liver Injury in Children. <i>Hepatology</i> , <b>2020</b> , 72, 1758-1770	11.2	27
98	A genome-wide association study of prostate cancer in Latinos. <i>International Journal of Cancer</i> , <b>2020</b> , 146, 1819-1826	7.5	13
97	A latent unknown clustering integrating multi-omics data (LUCID) with phenotypic traits. <i>Bioinformatics</i> , <b>2020</b> , 36, 842-850	7.2	9
96	Association of Fish Consumption and Mercury Exposure During Pregnancy With Metabolic Health and Inflammatory Biomarkers in Children. <i>JAMA Network Open</i> , <b>2020</b> , 3, e201007	10.4	10
95	Association of a Pathway-Specific Genetic Risk Score With Risk of Radiation-Associated Contralateral Breast Cancer. <i>JAMA Network Open</i> , <b>2019</b> , 2, e1912259	10.4	1
94	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , <b>2019</b> , 10, 431	17.4	45
93	DNA methylation patterns of adult survivors of adolescent/young adult Hodgkin lymphoma compared to their unaffected monozygotic twin. <i>Leukemia and Lymphoma</i> , <b>2019</b> , 60, 1429-1437	1.9	7
92	Reply to Xiaoling Lin, Brian T. Helfand, and Jianfeng Xu Letter to the Editor re: Daniel A. Leongamornlert, Edward J. Saunders, Sarah Wakerell, et al. Germline DNA Repair Gene Mutations in Young-onset Prostate Cancer Cases in the UK: Evidence for a More Extensive Genetic Panel. <i>Eur Urol</i> <b>2019</b> ;76:329-37. <i>European Urology</i> , <b>2019</b> , 76, e130-e131	10.2	
91	Elucidation of causal direction between asthma and obesity: a bi-directional Mendelian randomization study. <i>International Journal of Epidemiology</i> , <b>2019</b> , 48, 899-907	7.8	23

90	Perfluoroalkyl substances, metabolomic profiling, and alterations in glucose homeostasis among overweight and obese Hispanic children: A proof-of-concept analysis. <i>Environment International</i> , <b>2019</b> , 126, 445-453	12.9	54
89	Telomerase Variants in Patients with Cirrhosis Awaiting Liver Transplantation. <i>Hepatology</i> , <b>2019</b> , 69, 2652-2663	11.2	7
88	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. <i>Journal of the National Cancer Institute</i> , <b>2019</b> , 111, 146-157	9.7	67
87	Epstein-Barr virus load is higher in long-term Hodgkin lymphoma survivors compared to their unaffected twins and unrelated controls. <i>British Journal of Haematology</i> , <b>2019</b> , 185, 377-380	4.5	1
86	Genetic overlap between autoimmune diseases and non-Hodgkin lymphoma subtypes. <i>Genetic Epidemiology</i> , <b>2019</b> , 43, 844-863	2.6	15
85	Elevated numbers of PD-L1 expressing B cells are associated with the development of AIDS-NHL. <i>Scientific Reports</i> , <b>2019</b> , 9, 9371	4.9	12
84	Improvements to the Escalation with Overdose Control design and a comparison with the restricted Continual Reassessment Method. <i>Pharmaceutical Statistics</i> , <b>2019</b> , 18, 659-670	1	2
83	Genome-wide association study of circulating folate one-carbon metabolites. <i>Genetic Epidemiology</i> , <b>2019</b> , 43, 1030-1045	2.6	1
82	HiLDA: a statistical approach to investigate differences in mutational signatures. <i>PeerJ</i> , <b>2019</b> , 7, e7557	3.1	4
81	Comprehensive Investigation of White Blood Cell and Gene Expression Profiles As Risk Factors for Multiple Myeloma in African Americans. <i>Blood</i> , <b>2019</b> , 134, 4379-4379	2.2	
80	Large-Scale Genome-Wide Association Study of East Asians Identifies Loci Associated With Risk for Colorectal Cancer. <i>Gastroenterology</i> , <b>2019</b> , 156, 1455-1466	13.3	55
79	Germline DNA Repair Gene Mutations in Young-onset Prostate Cancer Cases in the UK: Evidence for a More Extensive Genetic Panel. <i>European Urology</i> , <b>2019</b> , 76, 329-337	10.2	28
78	IC-P-056: INTERACTION EFFECT OF APOE-4 AND SUBJECTIVE SLEEP QUALITY ON CORTICAL THICKNESS IN COGNITIVELY HEALTHY ADULTS <b>2019</b> , 15, P57-P57		
77	An admixture mapping meta-analysis implicates genetic variation at 18q21 with asthma susceptibility in Latinos. <i>Journal of Allergy and Clinical Immunology</i> , <b>2019</b> , 143, 957-969	11.5	20
76	Using Bayes model averaging to leverage both gene main effects and G × E interactions to identify genomic regions in genome-wide association studies. <i>Genetic Epidemiology</i> , <b>2019</b> , 43, 150-165	2.6	4
75	A Unified Model for the Analysis of Gene-Environment Interaction. <i>American Journal of Epidemiology</i> , <b>2019</b> , 188, 760-767	3.8	2
74	Circulating Metabolic Biomarkers of Screen-Detected Prostate Cancer in the ProtecT Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2019</b> , 28, 208-216	4	9
73	Discovery of common and rare genetic risk variants for colorectal cancer. <i>Nature Genetics</i> , <b>2019</b> , 51, 76-83	36.3	177

72	Genetic risk of prostate cancer in Ugandan men. <i>Prostate</i> , <b>2018</b> , 78, 370-376	4.2	17
71	Genome-wide association study and meta-analysis in Northern European populations replicate multiple colorectal cancer risk loci. <i>International Journal of Cancer</i> , <b>2018</b> , 142, 540-546	7.5	21
70	Rare germline variants in DNA repair genes and the angiogenesis pathway predispose prostate cancer patients to develop metastatic disease. <i>British Journal of Cancer</i> , <b>2018</b> , 119, 96-104	8.7	27
69	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
68	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. <i>Nature Genetics</i> , <b>2018</b> , 50, 928-936	36.3	34 <sup>o</sup>
67	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. <i>Nature Communications</i> , <b>2018</b> , 9, 2256	17.4	57
66	DNA repair gene panel mutations in young onset prostate cancer cases in the.. <i>Journal of Clinical Oncology</i> , <b>2018</b> , 36, 18-18	2.2	
65	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
64	Germline variation at 8q24 and prostate cancer risk in men of European ancestry. <i>Nature Communications</i> , <b>2018</b> , 9, 4616	17.4	30
63	Mendelian randomisation implicates hyperlipidaemia as a risk factor for colorectal cancer. <i>International Journal of Cancer</i> , <b>2017</b> , 140, 2701-2708	7.5	50
62	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
61	Novel colon cancer susceptibility variants identified from a genome-wide association study in African Americans. <i>International Journal of Cancer</i> , <b>2017</b> , 140, 2728-2733	7.5	17
60	Pro-inflammatory fatty acid profile and colorectal cancer risk: A Mendelian randomisation analysis. <i>European Journal of Cancer</i> , <b>2017</b> , 84, 228-238	7.5	56
59	Inherited variation in circadian rhythm genes and risks of prostate cancer and three other cancer sites in combined cancer consortia. <i>International Journal of Cancer</i> , <b>2017</b> , 141, 1794-1802	7.5	19
58	Quantifying the Genetic Correlation between Multiple Cancer Types. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2017</b> , 26, 1427-1435	4	25
57	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2017</b> , 26, 126-135	4	183
56	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
55	Discovery and fine-mapping of adiposity loci using high density imputation of genome-wide association studies in individuals of African ancestry: African Ancestry Anthropometry Genetics Consortium. <i>PLoS Genetics</i> , <b>2017</b> , 13, e1006719	6	60

54	Prostate cancer meta-analysis from more than 145,000 men to identify 65 novel prostate cancer susceptibility loci.. <i>Journal of Clinical Oncology</i> , <b>2017</b> , 2017, 1-1	2.2	4
53	Common variants in the obesity-associated genes FTO and MC4R are not associated with risk of colorectal cancer. <i>Cancer Epidemiology</i> , <b>2016</b> , 44, 1-4	2.8	9
52	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
51	Genome-wide association study of colorectal cancer in Hispanics. <i>Carcinogenesis</i> , <b>2016</b> , 37, 547-556	4.6	26
50	The contribution of rare variation to prostate cancer heritability. <i>Nature Genetics</i> , <b>2016</b> , 48, 30-5	36.3	106
49	Variation at 2q35 (PNKD and TMBIM1) influences colorectal cancer risk and identifies a pleiotropic effect with inflammatory bowel disease. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 2349-2359	5.6	27
48	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
47	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
46	HLA Haplotypes Are Associated with Multiple Myeloma Risk in the African American Multiple Myeloma Study (AAMMS). <i>Blood</i> , <b>2016</b> , 128, 3250-3250	2.2	1
45	Fine-Mapping of Common Genetic Variants Associated with Colorectal Tumor Risk Identified Potential Functional Variants. <i>PLoS ONE</i> , <b>2016</b> , 11, e0157521	3.7	5
44	Detecting Gene-Environment Interactions for a Quantitative Trait in a Genome-Wide Association Study. <i>Genetic Epidemiology</i> , <b>2016</b> , 40, 394-403	2.6	15
43	Mendelian randomisation analysis strongly implicates adiposity with risk of developing colorectal cancer. <i>British Journal of Cancer</i> , <b>2016</b> , 115, 266-72	8.7	39
42	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. <i>Cancer Research</i> , <b>2016</b> , 76, 5103-14	10.1	66
41	A Meta-analysis of Multiple Myeloma Risk Regions in African and European Ancestry Populations Identifies Putatively Functional Loci. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2016</b> , 25, 1609-1618	4	13
40	JAM: A Scalable Bayesian Framework for Joint Analysis of Marginal SNP Effects. <i>Genetic Epidemiology</i> , <b>2016</b> , 40, 188-201	2.6	41
39	UGT1A and UGT2B genetic variation alters nicotine and nitrosamine glucuronidation in european and african american smokers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2015</b> , 24, 94-104	4	24
38	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. <i>Nature Communications</i> , <b>2015</b> , 6, 7138	17.4	106
37	A new GWAS and meta-analysis with 1000Genomes imputation identifies novel risk variants for colorectal cancer. <i>Scientific Reports</i> , <b>2015</b> , 5, 10442	4.9	94

36	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
35	Identification of a Novel Mucin Gene HCG22 Associated With Steroid-Induced Ocular Hypertension <b>2015</b> , 56, 2737-48		22
34	Multiple novel prostate cancer susceptibility signals identified by fine-mapping of known risk loci among Europeans. <i>Human Molecular Genetics</i> , <b>2015</b> , 24, 5589-602	5.6	54
33	Identification of a common variant with potential pleiotropic effect on risk of inflammatory bowel disease and colorectal cancer. <i>Carcinogenesis</i> , <b>2015</b> , 36, 999-1007	4.6	21
32	Germline Mutations in the BRIP1, BARD1, PALB2, and NBN Genes in Women With Ovarian Cancer. <i>Journal of the National Cancer Institute</i> , <b>2015</b> , 107,	9.7	239
31	Enlight: web-based integration of GWAS results with biological annotations. <i>Bioinformatics</i> , <b>2015</b> , 31, 275-6	7.2	17
30	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. <i>Scientific Reports</i> , <b>2015</b> , 5, 17369	4.9	27
29	Drug Metabolizing Enzyme and Transporter Gene Variation, Nicotine Metabolism, Prospective Abstinence, and Cigarette Consumption. <i>PLoS ONE</i> , <b>2015</b> , 10, e0126113	3.7	14
28	Genetic ancestry influences asthma susceptibility and lung function among Latinos. <i>Journal of Allergy and Clinical Immunology</i> , <b>2015</b> , 135, 228-35	11.5	85
27	Cross-disorder genome-wide analyses suggest a complex genetic relationship between Tourette syndrome and OCD. <i>American Journal of Psychiatry</i> , <b>2015</b> , 172, 82-93	11.9	101
26	DNA Methylation Differences in Twins Discordant for Adolescent/Young Adult Hodgkin Lymphoma. <i>Blood</i> , <b>2015</b> , 126, 179-179	2.2	
25	Genome-wide interaction study of smoking and bladder cancer risk. <i>Carcinogenesis</i> , <b>2014</b> , 35, 1737-44	4.6	33
24	Identification and characterization of functional risk variants for colorectal cancer mapping to chromosome 11q23.1. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 2198-209	5.6	31
23	The 19q12 bladder cancer GWAS signal: association with cyclin E function and aggressive disease. <i>Cancer Research</i> , <b>2014</b> , 74, 5808-18	10.1	19
22	Identification of susceptibility loci for colorectal cancer in a genome-wide meta-analysis. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 4729-37	5.6	107
21	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
20	A novel colorectal cancer risk locus at 4q32.2 identified from an international genome-wide association study. <i>Carcinogenesis</i> , <b>2014</b> , 35, 2512-9	4.6	25
19	Pleiotropic effects of genetic risk variants for other cancers on colorectal cancer risk: PAGE, GECCO and CCFR consortia. <i>Gut</i> , <b>2014</b> , 63, 800-7	19.2	27



18	Organic cation transporter variation and response to smoking cessation therapies. <i>Nicotine and Tobacco Research</i> , <b>2014</b> , 16, 1638-46	4.9	18
17	Nicotine dependence as a moderator of genetic influences on smoking cessation treatment outcome. <i>Drug and Alcohol Dependence</i> , <b>2014</b> , 138, 109-17	4.9	12
16	Meta-Analysis of Hodgkin Lymphoma and Asthma Genome-Wide Association Scans reveals common variants in GATA3. <i>Blood</i> , <b>2014</b> , 124, 135-135	2.2	
15	Genetic Susceptibility Markers of Multiple Myeloma in African-Americans. <i>Blood</i> , <b>2014</b> , 124, 2030-2030	2.2	
14	Identification of Genetic Susceptibility Loci for Colorectal Tumors in a Genome-Wide Meta-analysis. <i>Gastroenterology</i> , <b>2013</b> , 144, 799-807.e24	13.3	250
13	Partitioning the heritability of Tourette syndrome and obsessive compulsive disorder reveals differences in genetic architecture. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003864	6	189
12	Sneaker "jack" males outcompete dominant "hooknose" males under sperm competition in Chinook salmon ( <i>Oncorhynchus tshawytscha</i> ). <i>Ecology and Evolution</i> , <b>2013</b> , 3, 4987-97	2.8	24
11	Meta-analysis of new genome-wide association studies of colorectal cancer risk. <i>Human Genetics</i> , <b>2012</b> , 131, 217-34	6.3	173
10	Joint analysis for integrating two related studies of different data types and different study designs using hierarchical modeling approaches. <i>Human Heredity</i> , <b>2012</b> , 74, 83-96	1.1	5
9	Genome-wide search for gene-gene interactions in colorectal cancer. <i>PLoS ONE</i> , <b>2012</b> , 7, e52535	3.7	29
8	Genotype-environment interactions in microsatellite stable/microsatellite instability-low colorectal cancer: results from a genome-wide association study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2011</b> , 20, 758-66	4	47
7	Discovery of complex pathways from observational data. <i>Statistics in Medicine</i> , <b>2010</b> , 29, 1998-2011	2.3	25
6	Use of pathway information in molecular epidemiology. <i>Human Genomics</i> , <b>2009</b> , 4, 21-42	6.8	38
5	Nicotinic acetylcholine receptor beta2 subunit gene implicated in a systems-based candidate gene study of smoking cessation. <i>Human Molecular Genetics</i> , <b>2008</b> , 17, 2834-48	5.6	118
4	Multistage Genetic Association Studies <b>2007</b> , 1		4
3	A testing framework for identifying susceptibility genes in the presence of epistasis. <i>American Journal of Human Genetics</i> , <b>2006</b> , 78, 15-27	11	154
2	SNPs, haplotypes, and model selection in a candidate gene region: the SIMPLE analysis for multilocus data. <i>Genetic Epidemiology</i> , <b>2004</b> , 27, 429-41	2.6	33
1	Bayesian modeling of complex metabolic pathways. <i>Human Heredity</i> , <b>2003</b> , 56, 83-93	1.1	78

