David V Conti

List of Publications by Year
in descending order

157 papers	8,200 citations	70961 41 h-index	62479 80 g-index
171 all docs	171 docs citations	171 times ranked	13368 citing authors

#	Article	IF	Citations
1	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. Nature Genetics, 2018, 50, 928-936.	9.4	652
2	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. Nature Genetics, 2014, 46, 1103-1109.	9.4	408
3	Discovery of common and rare genetic risk variants for colorectal cancer. Nature Genetics, 2019, 51, 76-87.	9.4	377
4	Germline Mutations in the BRIP1, BARD1, PALB2, and NBN Genes in Women With Ovarian Cancer. Journal of the National Cancer Institute, 2015, 107, .	3.0	311
5	Identification of Genetic Susceptibility Loci for Colorectal Tumors in a Genome-Wide Meta-analysis. Gastroenterology, 2013, 144, 799-807.e24.	0.6	292
6	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 126-135.	1.1	278
7	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. Nature Genetics, 2021, 53, 65-75.	9.4	264
8	Partitioning the Heritability of Tourette Syndrome and Obsessive Compulsive Disorder Reveals Differences in Genetic Architecture. PLoS Genetics, 2013, 9, e1003864.	1.5	241
9	Meta-analysis of new genome-wide association studies of colorectal cancer risk. Human Genetics, 2012, 131, 217-234.	1.8	183
10	A Testing Framework for Identifying Susceptibility Genes in the Presence of Epistasis. American Journal of Human Genetics, 2006, 78, 15-27.	2.6	174
11	The contribution of rare variation to prostate cancer heritability. Nature Genetics, 2016, 48, 30-35.	9.4	139
12	Genome-wide association study of colorectal cancer identifies six new susceptibility loci. Nature Communications, 2015, 6, 7138.	5.8	138
13	Nicotinic acetylcholine receptor Î²2 subunit gene implicated in a systems-based candidate gene study of smoking cessation. Human Molecular Genetics, 2008, 17, 2834-2848.	1.4	129
14	Novel Common Genetic Susceptibility Loci for Colorectal Cancer. Journal of the National Cancer Institute, 2019, 111, 146-157.	3.0	129
15	Identification of susceptibility loci for colorectal cancer in a genome-wide meta-analysis. Human Molecular Genetics, 2014, 23, 4729-4737.	1.4	128
16	Exposure to per- and Polyfluoroalkyl Substances and Markers of Liver Injury: A Systematic Review and Meta-Analysis. Environmental Health Perspectives, 2022, 130, 46001.	2.8	128
17	Cross-Disorder Genome-Wide Analyses Suggest a Complex Genetic Relationship Between Tourette's Syndrome and OCD. American Journal of Psychiatry, 2015, 172, 82-93.	4.0	117
18	Genetic ancestry influences asthma susceptibility and lung function among Latinos. Journal of Allergy and Clinical Immunology, 2015, 135, 228-235.	1.5	113

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19	Prostate Cancer Susceptibility in Men of African Ancestry at 8q24. Journal of the National Cancer Institute, 2016, 108, djv431.	3.0	111
20	Large-Scale Genome-Wide Association Study of East Asians Identifies Loci Associated With Risk for Colorectal Cancer. Gastroenterology, 2019, 156, 1455-1466.	0.6	111
21	Cumulative Burden of Colorectal Cancerâ€"Associated Genetic Variants Is More Strongly Associated With Early-Onset vs Late-Onset Cancer. Gastroenterology, 2020, 158, 1274-1286.e12.	0.6	110
22	Perfluoroalkyl substances and severity of nonalcoholic fatty liver in Children: An untargeted metabolomics approach. Environment International, 2020, 134, 105220.	4.8	110
23	A new GWAS and meta-analysis with 1000Genomes imputation identifies novel risk variants for colorectal cancer. Scientific Reports, 2015, 5, 10442.	1.6	109
24	Perfluoroalkyl substances, metabolomic profiling, and alterations in glucose homeostasis among overweight and obese Hispanic children: A proof-of-concept analysis. Environment International, 2019, 126, 445-453.	4.8	105
25	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. Cancer Research, 2016, 76, 5103-5114.	0.4	100
26	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. PLoS Genetics, 2017, 13, e1006719.	1.5	98
27	Prenatal Exposure to Perfluoroalkyl Substances Associated With Increased Susceptibility to Liver Injury in Children. Hepatology, 2020, 72, 1758-1770.	3.6	90
28	Bayesian Modeling of Complex Metabolic Pathways. Human Heredity, 2003, 56, 83-93.	0.4	88
29	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. Nature Communications, 2018, 9, 2256.	5.8	88
30	Shared heritability and functional enrichment across six solid cancers. Nature Communications, 2019, 10, 431.	5.8	88
31	Pro-inflammatory fatty acid profile and colorectal cancer risk: A Mendelian randomisation analysis. European Journal of Cancer, 2017, 84, 228-238.	1.3	81
32	Mendelian randomisation implicates hyperlipidaemia as a risk factor for colorectal cancer. International Journal of Cancer, 2017, 140, 2701-2708.	2.3	76
33	JAM: A Scalable Bayesian Framework for Joint Analysis of Marginal SNP Effects. Genetic Epidemiology, 2016, 40, 188-201.	0.6	74
34	Multiple novel prostate cancer susceptibility signals identified by fine-mapping of known risk loci among Europeans. Human Molecular Genetics, 2015, 24, 5589-5602.	1.4	67
35	Mendelian randomisation analysis strongly implicates adiposity with risk of developing colorectal cancer. British Journal of Cancer, 2016, 115, 266-272.	2.9	57
36	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. Journal of the National Cancer Institute, 2017, 109, .	3.0	57

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37	Genotype–Environment Interactions in Microsatellite Stable/Microsatellite Instability-Low Colorectal Cancer: Results from a Genome-Wide Association Study. Cancer Epidemiology Biomarkers and Prevention, 2011, 20, 758-766.	1.1	50
38	Genome-wide interaction study of smoking and bladder cancer risk. Carcinogenesis, 2014, 35, 1737-1744.	1.3	50
39	Integration of multiethnic fine-mapping and genomic annotation to prioritize candidate functional SNPs at prostate cancer susceptibility regions. Human Molecular Genetics, 2015, 24, 5603-5618.	1.4	50
40	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. Nature Communications, 2016, 7, 10979.	5.8	50
41	Quantifying the Genetic Correlation between Multiple Cancer Types. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 1427-1435.	1.1	48
42	Germline DNA Repair Gene Mutations in Young-onset Prostate Cancer Cases in the UK: Evidence for a More Extensive Genetic Panel. European Urology, 2019, 76, 329-337.	0.9	48
43	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.	0.8	44
44	Genetic architectures of proximal and distal colorectal cancer are partly distinct. Gut, 2021, 70, 1325-1334.	6.1	44
45	Germline variation at 8q24 and prostate cancer risk in men of European ancestry. Nature Communications, 2018, 9, 4616.	5.8	43
46	Use of pathway information in molecular epidemiology. Human Genomics, 2009, 4, 21.	1.4	42
47	SNPs, haplotypes, and model selection in a candidate gene region: The SIMPle analysis for multilocus data. Genetic Epidemiology, 2004, 27, 429-441.	0.6	40
48	Rare germline variants in DNA repair genes and the angiogenesis pathway predispose prostate cancer patients to develop metastatic disease. British Journal of Cancer, 2018, 119, 96-104.	2.9	40
49	Germline Sequencing DNA Repair Genes in 5545 Men With Aggressive and Nonaggressive Prostate Cancer. Journal of the National Cancer Institute, 2021, 113, 616-625.	3.0	40
50	Evaluation of a Multiethnic Polygenic Risk Score Model for Prostate Cancer. Journal of the National Cancer Institute, 2022, 114, 771-774.	3.0	39
51	Combined Effect of a Polygenic Risk Score and Rare Genetic Variants on Prostate Cancer Risk. European Urology, 2021, 80, 134-138.	0.9	39
52	Variation at 2q35 (*PNKD* and *TMBIM1*) influences colorectal cancer risk and identifies a pleiotropic effect with inflammatory bowel disease. Human Molecular Genetics, 2016, 25, 2349-2359.	1.4	37
53	Elucidation of causal direction between asthma and obesity: a bi-directional Mendelian randomization study. International Journal of Epidemiology, 2019, 48, 899-907.	0.9	37
54	Identification and characterization of functional risk variants for colorectal cancer mapping to chromosome 11q23.1. Human Molecular Genetics, 2014, 23, 2198-2209.	1.4	36

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55	Identifying Novel Susceptibility Genes for Colorectal Cancer Risk From a Transcriptome-Wide Association Study of 125,478 Subjects. Gastroenterology, 2021, 160, 1164-1178.e6.	0.6	36
56	Pleiotropic effects of genetic risk variants for other cancers on colorectal cancer risk: PAGE, GECCO and CCFR consortia. Gut, 2014, 63, 800-807.	6.1	35
57	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. Scientific Reports, 2015, 5, 17369.	1.6	35
58	Genome-Wide Search for Gene-Gene Interactions in Colorectal Cancer. PLoS ONE, 2012, 7, e52535.	1.1	35
59	Detecting Geneâ€"Environment Interactions for a Quantitative Trait in a Genomeâ€Wide Association Study. Genetic Epidemiology, 2016, 40, 394-403.	0.6	34
60	Genome-wide association study of colorectal cancer in Hispanics. Carcinogenesis, 2016, 37, 547-556.	1.3	34
61	A Cross-Sectional Study Examining the Seroprevalence of Severe Acute Respiratory Syndrome Coronavirus 2 Antibodies in a University Student Population. Journal of Adolescent Health, 2020, 67, 763-768.	1.2	34
62	Prenatal metal mixtures and child blood pressure in the Rhea mother-child cohort in Greece. Environmental Health, 2021, 20, 1.	1.7	34
63	Sneaker â€œjackâ€males outcompete dominant â€œhooknoseâ€males under sperm competition in C hinook salmon (O ncorhynchus tshawytscha). Ecology and Evolution, 2013, 3, 4987-4997.	0.8	33
64	An admixture mapping meta-analysis implicates genetic variation at 18q21 with asthma susceptibility in Latinos. Journal of Allergy and Clinical Immunology, 2019, 143, 957-969.	1.5	33
65	Prenatal and postnatal exposure to PFAS and cardiometabolic factors and inflammation status in children from six European cohorts. Environment International, 2021, 157, 106853.	4.8	33
66	A Germline Variant at 8q24 Contributes to Familial Clustering of Prostate Cancer in Men of African Ancestry. European Urology, 2020, 78, 316-320.	0.9	32
67	Genetic risk of prostate cancer in Ugandan men. Prostate, 2018, 78, 370-376.	1.2	31
68	Prenatal exposure to persistent organic pollutants and childhood obesity: A systematic review and metaâ€analysis of human studies. Obesity Reviews, 2022, 23, e13383.	3.1	31
69	A novel colorectal cancer risk locus at 4q32.2 identified from an international genome-wide association study. Carcinogenesis, 2014, 35, 2512-2519.	1.3	30
70	Pathogenic Variants in Cancer Predisposition Genes and Prostate Cancer Risk in Men of African Ancestry. JCO Precision Oncology, 2020, 4, 32-43.	1.5	30
71	Association of Fish Consumption and Mercury Exposure During Pregnancy With Metabolic Health and Inflammatory Biomarkers in Children. JAMA Network Open, 2020, 3, e201007.	2.8	30
72	Association of Prenatal Exposure to Endocrine-Disrupting Chemicals With Liver Injury in Children. JAMA Network Open, 2022, 5, e2220176.	2.8	30

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73	Identification of a Novel Mucin Gene *HCG22* Associated With Steroid-Induced Ocular Hypertension. , 2015, 56, 2737.		28
74	Identification of a common variant with potential pleiotropic effect on risk of inflammatory bowel disease and colorectal cancer. Carcinogenesis, 2015, 36, 999-1007.	1.3	28
75	Inherited variation in circadian rhythm genes and risks of prostate cancer and three other cancer sites in combined cancer consortia. International Journal of Cancer, 2017, 141, 1794-1802.	2.3	28
76	Genetic overlap between autoimmune diseases and non‑Hodgkin lymphoma subtypes. Genetic Epidemiology, 2019, 43, 844-863.	0.6	28
77	An integrative multi-omics analysis to identify candidate DNA methylation biomarkers related to prostate cancer risk. Nature Communications, 2020, 11, 3905.	5.8	28
78	Discovery of complex pathways from observational data. Statistics in Medicine, 2010, 29, 1998-2011.	0.8	27
79	*UGT1A* and *UGT2B* Genetic Variation Alters Nicotine and Nitrosamine Glucuronidation in European and African American Smokers. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 94-104.	1.1	27
80	Whole-exome sequencing of over 4100 men of African ancestry and prostate cancer risk. Human Molecular Genetics, 2016, 25, 371-381.	1.4	26
81	Novel colon cancer susceptibility variants identified from a genome‑wide association study in African Americans. International Journal of Cancer, 2017, 140, 2728-2733.	2.3	26
82	Genome‑wide association study and meta‑analysis in Northern European populations replicate multiple colorectal cancer risk loci. International Journal of Cancer, 2018, 142, 540-546.	2.3	26
83	Mendelian Randomization of Circulating Polyunsaturated Fatty Acids and Colorectal Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 860-870.	1.1	26
84	The 19q12 Bladder Cancer GWAS Signal: Association with Cyclin E Function and Aggressive Disease. Cancer Research, 2014, 74, 5808-5818.	0.4	24
85	Characterizing Genetic Susceptibility to Breast Cancer in Women of African Ancestry. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 1016-1026.	1.1	24
86	A genome‑wide association study of prostate cancer in Latinos. International Journal of Cancer, 2020, 146, 1819-1826.	2.3	24
87	Cross-ancestry GWAS meta-analysis identifies six breast cancer loci in African and European ancestry women. Nature Communications, 2021, 12, 4198.	5.8	24
88	Advancing tools for human early lifecourse exposome research and translation (ATHLETE). Environmental Epidemiology, 2021, 5, e166.	1.4	24
89	The Four-Kallikrein Panel Is Effective in Identifying Aggressive Prostate Cancer in a Multiethnic Population. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1381-1388.	1.1	22
90	Association of Genetic Risk Score With NAFLD in An Ethnically Diverse Cohort. Hepatology Communications, 2021, 5, 1689-1703.	2.0	22

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91	In Utero Exposure to Mercury Is Associated With Increased Susceptibility to Liver Injury and Inflammation in Childhood. Hepatology, 2021, 74, 1546-1559.	3.6	22
92	A Rare Germline HOXB13 Variant Contributes to Risk of Prostate Cancer in Men of African Ancestry. European Urology, 2022, 81, 458-462.	0.9	22
93	Organic Cation Transporter Variation and Response to Smoking Cessation Therapies. Nicotine and Tobacco Research, 2014, 16, 1638-1646.	1.4	21
94	Circulating Metabolic Biomarkers of Screen-Detected Prostate Cancer in the ProtecT Study. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 208-216.	1.1	21
95	Drug Metabolizing Enzyme and Transporter Gene Variation, Nicotine Metabolism, Prospective Abstinence, and Cigarette Consumption. PLoS ONE, 2015, 10, e0126113.	1.1	20
96	Elevated numbers of PD-L1 expressing B cells are associated with the development of AIDS-NHL. Scientific Reports, 2019, 9, 9371.	1.6	19
97	Exposure to Perfluoroalkyl Substances and Glucose Homeostasis in Youth. Environmental Health Perspectives, 2021, 129, 97002.	2.8	19
98	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.	1.1	18
99	A latent unknown clustering integrating multi-omics data (LUCID) with phenotypic traits. Bioinformatics, 2020, 36, 842-850.	1.8	18
100	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. American Journal of Human Genetics, 2021, 108, 564-582.	2.6	18
101	Enlight: web-based integration of GWAS results with biological annotations. Bioinformatics, 2015, 31, 275-276.	1.8	17
102	Environmental chemical burden in metabolic tissues and systemic biological pathways in adolescent bariatric surgery patients: A pilot untargeted metabolomic approach. Environment International, 2020, 143, 105957.	4.8	17
103	A meta-analysis of genome-wide association studies of multiple myeloma among men and women of African ancestry. Blood Advances, 2020, 4, 181-190.	2.5	16
104	Circulating insulin-like growth factors and risks of overall, aggressive and early-onset prostate cancer: a collaborative analysis of 20 prospective studies and Mendelian randomization analysis. International Journal of Epidemiology, 2023, 52, 71-86.	0.9	16
105	A Unified Model for the Analysis of Gene-Environment Interaction. American Journal of Epidemiology, 2019, 188, 760-767.	1.6	15
106	Validation of a multi-ancestry polygenic risk score and age-specific risks of prostate cancer: A meta-analysis within diverse populations. ELife, 0, 11, .	2.8	15
107	Nicotine dependence as a moderator of genetic influences on smoking cessation treatment outcome. Drug and Alcohol Dependence, 2014, 138, 109-117.	1.6	13
108	Evaluation of 71 Coronary Artery Disease Risk Variants in a Multiethnic Cohort. Frontiers in Cardiovascular Medicine, 2018, 5, 19.	1.1	13

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109	Common variants in the obesity-associated genes FTO and MC4R are not associated with risk of colorectal cancer. Cancer Epidemiology, 2016, 44, 1-4.	0.8	12
110	DNA methylation patterns of adult survivors of adolescent/young adult Hodgkin lymphoma compared to their unaffected monozygotic twin. Leukemia and Lymphoma, 2019, 60, 1429-1437.	0.6	11
111	Interaction effect of alcohol consumption and Alzheimer disease polygenic risk score on the brain cortical thickness of cognitively normal subjects. Alcohol, 2020, 85, 1-12.	0.8	11
112	Replication and Genetic Risk Score Analysis for Pancreatic Cancer in a Diverse Multiethnic Population. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 2686-2692.	1.1	11
113	Polygenic risk scores for prediction of breast cancer risk in women of African ancestry: a cross-ancestry approach. Human Molecular Genetics, 2022, 31, 3133-3143.	1.4	11
114	Telomerase Variants in Patients with Cirrhosis Awaiting Liver Transplantation. Hepatology, 2019, 69, 2652-2663.	3.6	10
115	Bayesian variable selection with a pleiotropic loss function in Mendelian randomization. Statistics in Medicine, 2021, 40, 5025-5045.	0.8	9
116	Plasma concentrations of lipophilic persistent organic pollutants and glucose homeostasis in youth populations. Environmental Research, 2022, 212, 113296.	3.7	9
117	Fine-Mapping of Common Genetic Variants Associated with Colorectal Tumor Risk Identified Potential Functional Variants. PLoS ONE, 2016, 11, e0157521.	1.1	8
118	Variability in Cytogenetic Testing for Multiple Myeloma: A Comprehensive Analysis From Across the United States. JCO Oncology Practice, 2020, 16, e1169-e1180.	1.4	8
119	Functional informed genomeâ€wide interaction analysis of body mass index, diabetes and colorectal cancer risk. Cancer Medicine, 2020, 9, 3563-3573.	1.3	7
120	Novel strategy for disease risk prediction incorporating predicted gene expression and DNA methylation data: a multiâ€phased study of prostate cancer. Cancer Communications, 2021, 41, 1387-1397.	3.7	6
121	Urinary metabolic biomarkers of diet quality in European children are associated with metabolic health. ELife, 2022, 11, .	2.8	6
122	Beyond GWAS of Colorectal Cancer: Evidence of Interaction with Alcohol Consumption and Putative Causal Variant for the 10q24.2 Region. Cancer Epidemiology Biomarkers and Prevention, 2022, 31, 1077-1089.	1.1	6
123	Characteristics associated with COVID-19 vaccination status among staff and faculty of a large, diverse University in Los Angeles: The Trojan Pandemic Response Initiative. Preventive Medicine Reports, 2022, 27, 101802.	0.8	6
124	Joint Analysis for Integrating Two Related Studies of Different Data Types and Different Study Designs Using Hierarchical Modeling Approaches. Human Heredity, 2012, 74, 83-96.	0.4	5
125	Association of a Pathway-Specific Genetic Risk Score With Risk of Radiation-Associated Contralateral Breast Cancer. JAMA Network Open, 2019, 2, e1912259.	2.8	5
126	Prenatal and childhood exposure to air pollution and traffic and the risk of liver injury in European children. Environmental Epidemiology, 2021, 5, e153.	1.4	5

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127	An integrated risk and epidemiological model to estimate risk-stratified COVID-19 outcomes for Los Angeles County: March 1, 2020â€"March 1, 2021. PLoS ONE, 2021, 16, e0253549.	1.1	5
128	HiLDA: a statistical approach to investigate differences in mutational signatures. PeerJ, 2019, 7, e7557.	0.9	5
129	Prostate cancer meta-analysis from more than 145,000 men to identify 65 novel prostate cancer susceptibility loci.. Journal of Clinical Oncology, 2017, 2017, 1-1.	0.8	5
130	Using Bayes model averaging to leverage both gene main effects and Gâ€‰Ã—â€‰E interactions to identify genomic regions in genomeâ€wide association studies. Genetic Epidemiology, 2019, 43, 150-165.	0.6	4
131	Infectious mononucleosis, immune genotypes, and non-Hodgkin lymphoma (NHL): an InterLymph Consortium study. Cancer Causes and Control, 2020, 31, 451-462.	0.8	4
132	A Hierarchical Approach Using Marginal Summary Statistics for Multiple Intermediates in a Mendelian Randomization or Transcriptome Analysis. American Journal of Epidemiology, 2021, 190, 1148-1158.	1.6	3
133	Rare Variants in the DNA Repair Pathway and the Risk of Colorectal Cancer. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 895-903.	1.1	3
134	Ancestral diversity improves discovery and fine-mapping of genetic loci for anthropometric traitsâ€"The Hispanic/Latino Anthropometry Consortium. Human Genetics and Genomics Advances, 2022, 3, 100099.	1.0	3
135	Improvements to the Escalation with Overdose Control design and a comparison with the restricted Continual Reassessment Method. Pharmaceutical Statistics, 2019, 18, 659-670.	0.7	2
136	Genomeâ€wide association study of circulating folate oneâ€carbon metabolites. Genetic Epidemiology, 2019, 43, 1030-1045.	0.6	2
137	Lymphoma-Associated Biomarkers Are Increased in Current Smokers in Twin Pairs Discordant for Smoking. Cancers, 2021, 13, 5395.	1.7	2
138	Epsteinâ€"Barr virus load is higher in longâ€term Hodgkin lymphoma survivors compared to their unaffected twins and unrelated controls. British Journal of Haematology, 2019, 185, 377-380.	1.2	1
139	Sign-based Shrinkage Based on an Asymmetric LASSO Penalty. Journal of Data Science, 2021, 19, 429-449.	0.5	1
140	Exposure to perfluoroalkyl substances (PFAS) and liver injury: a systematic review and meta-analysis. ISEE Conference Abstracts, 2021, 2021, .	0.0	1
141	HLA Haplotypes Are Associated with Multiple Myeloma Risk in the African American Multiple Myeloma Study (AAMMS). Blood, 2016, 128, 3250-3250.	0.6	1
142	Meta-Analysis of Hodgkin Lymphoma and Asthma Genome-Wide Association Scans reveals common variants in GATA3. Blood, 2014, 124, 135-135.	0.6	1
143	DNA Methylation Differences in Twins Discordant for Adolescent/Young Adult Hodgkin Lymphoma. Blood, 2015, 126, 179-179.	0.6	1
144	Whole-Exome Sequencing in Multiplex Families to Identify Novel AYA Classical Hodgkin Lymphoma Predisposition Genes. Blood, 2021, 138, 3499-3499.	0.6	1

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145	Reply to Xiaoling Lin, Brian T. Helfand, and Jianfeng Xu's 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. Eur Urol 2019;76:329–37. European Urology, 2019, 76, e130-e131.	0.9	0
146	ICâ€Pâ€056: INTERACTION EFFECT OF APOEâ€4 AND SUBJECTIVE SLEEP QUALITY ON CORTICAL THICKNESS IN COGNITIVELY HEALTHY ADULTS. Alzheimer's and Dementia, 2019, 15, P57.	0.4	0
147	Abstract LB011: Meta-analysis in more than 80,000 men of African ancestry identified nine novel variants associated with prostate cancer. , 2021, , .		0
148	Exposure to lipophilic chemicals and glucose homeostasis in youth. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
149	Having your cake (mix) and eating it too: Independent, interaction, and group effects of mixtures using Bayesian Hierarchical Regression Modelling. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
150	Prenatal Metal Mixtures and Child Blood Pressure in the Rhea Mother-Child Cohort. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
151	Associations between liver PFAS concentrations and plasma extracellular miRNAs in a cohort of adolescents undergoing bariatric surgery. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
152	Genetic Susceptibility Markers of Multiple Myeloma in African-Americans. Blood, 2014, 124, 2030-2030.	0.6	0
153	DNA repair gene panel mutations in young onset prostate cancer cases in the.. Journal of Clinical Oncology, 2018, 36, 18-18.	0.8	0
154	Comprehensive Investigation of White Blood Cell and Gene Expression Profiles As Risk Factors for Multiple Myeloma in African Americans. Blood, 2019, 134, 4379-4379.	0.6	0
155	Differential Gene Expression in Circulating T-Cells in Long-Term Adolescent/Young Adult Hodgkin Lymphoma (AYAHL) Survivors and Their Unaffected Twins. Blood, 2021, 138, 1332-1332.	0.6	0
156	Abstract PR-16: Serum prostate-specific antigen levels and prostate cancer risk in a multiethnic population. , 2022, , .		0
157	OUP accepted manuscript. Journal of the National Cancer Institute, 2022, , .	3.0	0