

# Elizabeth R Hauser

## List of Publications by Year in descending order

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144  
papers

6,199  
citations

66343

42  
h-index

85541

71  
g-index

154  
all docs

154  
docs citations

154  
times ranked

9264  
citing authors

#	ARTICLE	IF	CITATIONS
1	Discovery of 318 new risk loci for type 2 diabetes and related vascular outcomes among 1.4 million participants in a multi-ancestry meta-analysis. <i>Nature Genetics</i> , 2020, 52, 680-691.	21.4	445
2	Association of a Peripheral Blood Metabolic Profile With Coronary Artery Disease and Risk of Subsequent Cardiovascular Events. <i>Circulation: Cardiovascular Genetics</i> , 2010, 3, 207-214.	5.1	390
3	Baseline metabolomic profiles predict cardiovascular events in patients at risk for coronary artery disease. <i>American Heart Journal</i> , 2012, 163, 844-850.e1.	2.7	271
4	Affected-sib-pair interval mapping and exclusion for complex genetic traits: Sampling considerations. , 1996, 13, 117-137.		198
5	The Finlandâ€“United States Investigation of Nonâ€“Insulinâ€“Dependent Diabetes Mellitus Genetics (FUSION) Study. I. An Autosomal Genome Scan for Genes That Predispose to Type 2 Diabetes. <i>American Journal of Human Genetics</i> , 2000, 67, 1174-1185.	6.2	186
6	Harmonizing Genetic Ancestry and Self-identified Race/Ethnicity in Genome-wide Association Studies. <i>American Journal of Human Genetics</i> , 2019, 105, 763-772.	6.2	169
7	Ordered subset analysis in genetic linkage mapping of complex traits. <i>Genetic Epidemiology</i> , 2004, 27, 53-63.	1.3	154
8	A Genomewide Scan for Early-Onset Coronary Artery Disease in 438 Families: The GENECARD Study. <i>American Journal of Human Genetics</i> , 2004, 75, 436-447.	6.2	152
9	Novel loci and pathways significantly associated with longevity. <i>Scientific Reports</i> , 2016, 6, 21243.	3.3	145
10	Prospective treatment of urea cycle disorders. <i>Journal of Pediatrics</i> , 1991, 119, 923-928.	1.8	143
11	High heritability of metabolomic profiles in families burdened with premature cardiovascular disease. <i>Molecular Systems Biology</i> , 2009, 5, 258.	7.2	140
12	The Finlandâ€“United States Investigation of Nonâ€“Insulin-Dependent Diabetes Mellitus Genetics (FUSION) Study. II. An Autosomal Genome Scan for Diabetes-Related Quantitative-Trait Loci. <i>American Journal of Human Genetics</i> , 2000, 67, 1186-1200.	6.2	121
13	Genotyping Array Design and Data Quality Control in the Million Veteran Program. <i>American Journal of Human Genetics</i> , 2020, 106, 535-548.	6.2	118
14	Familiality of Quantitative Metabolic Traits in Finnish Families with Non-Insulin-Dependent Diabetes mellitus. <i>Human Heredity</i> , 1999, 49, 159-168.	0.8	115
15	Dissecting the Shared Genetic Architecture of Suicide Attempt, Psychiatric Disorders, and Known Risk Factors. <i>Biological Psychiatry</i> , 2022, 91, 313-327.	1.3	114
16	Genome-wide analysis identifies novel susceptibility loci for myocardial infarction. <i>European Heart Journal</i> , 2021, 42, 919-933.	2.2	113
17	Peakwide Mapping on Chromosome 3q13 Identifies the Kalirin Gene as a Novel Candidate Gene for Coronary Artery Disease. <i>American Journal of Human Genetics</i> , 2007, 80, 650-663.	6.2	110
18	Validation of the association between a branched chain amino acid metabolite profile and extremes of coronary artery disease in patients referred for cardiac catheterization. <i>Atherosclerosis</i> , 2014, 232, 191-196.	0.8	109

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19	Ordered-Subsets Linkage Analysis Detects Novel Alzheimer Disease Loci on Chromosomes 2q34 and 15q22. <i>American Journal of Human Genetics</i> , 2003, 73, 1041-1051.	6.2	99
20	Accounting for Linkage in Family-Based Tests of Association with Missing Parental Genotypes. <i>American Journal of Human Genetics</i> , 2003, 73, 1016-1026.	6.2	89
21	Neuropeptide Y Gene Polymorphisms Confer Risk of Early-Onset Atherosclerosis. <i>PLoS Genetics</i> , 2009, 5, e1000318.	3.5	87
22	Early Adult-Onset POAG Linked to 15q11-13 Using Ordered Subset Analysis. , 2005, 46, 2002.		86
23	Association of long-term PM2.5 exposure with traditional and novel lipid measures related to cardiovascular disease risk. <i>Environment International</i> , 2019, 122, 193-200.	10.0	83
24	GATA2 Is Associated with Familial Early-Onset Coronary Artery Disease. <i>PLoS Genetics</i> , 2006, 2, e139.	3.5	82
25	Metabolomic Quantitative Trait Loci (mQTL) Mapping Implicates the Ubiquitin Proteasome System in Cardiovascular Disease Pathogenesis. <i>PLoS Genetics</i> , 2015, 11, e1005553.	3.5	81
26	Linkage Disequilibrium Inflates Type I Error Rates in Multipoint Linkage Analysis when Parental Genotypes Are Missing. <i>Human Heredity</i> , 2005, 59, 220-227.	0.8	74
27	A Large Set of Finnish Affected Sibling Pair Families With Type 2 Diabetes Suggests Susceptibility Loci on Chromosomes 6, 11, and 14. <i>Diabetes</i> , 2004, 53, 821-829.	0.6	73
28	The Finlandâ€“United States Investigation of Nonâ€“Insulin-Dependent Diabetes Mellitus Genetics (FUSION) Study. I. An Autosomal Genome Scan for Genes That Predispose to Type 2 Diabetes. <i>American Journal of Human Genetics</i> , 2000, 67, 1174-1185.	6.2	71
29	Association between satellite-based estimates of long-term PM2.5 exposure and coronary artery disease. <i>Environmental Research</i> , 2016, 145, 9-17.	7.5	69
30	Ozone exposure is associated with acute changes in inflammation, fibrinolysis, and endothelial cell function in coronary artery disease patients. <i>Environmental Health</i> , 2017, 16, 126.	4.0	67
31	Comprehensive genetic analysis of the platelet activating factor acetylhydrolase (PLA2G7) gene and cardiovascular disease in caseâ€“control and family datasets. <i>Human Molecular Genetics</i> , 2008, 17, 1318-1328.	2.9	66
32	A Guide for a Cardiovascular Genomics Biorepository: the CATHGEN Experience. <i>Journal of Cardiovascular Translational Research</i> , 2015, 8, 449-457.	2.4	64
33	Fine particulate matter and cardiovascular disease: Comparison of assessment methods for long-term exposure. <i>Environmental Research</i> , 2017, 159, 16-23.	7.5	63
34	Association of maternal IL-1 receptor antagonist intron 2 gene polymorphism and preterm birth. <i>American Journal of Obstetrics and Gynecology</i> , 2006, 195, 1249-1253.	1.3	57
35	Genetic Linkage Analysis of Complex Genetic Traits by Using Affected Sibling Pairs. <i>Biometrics</i> , 1998, 54, 1238.	1.4	55
36	Design of the Genetics of Early Onset Cardiovascular Disease (GENECARD) study. <i>American Heart Journal</i> , 2003, 145, 602-613.	2.7	55

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37	Genetic effects in the leukotriene biosynthesis pathway and association with atherosclerosis. <i>Human Genetics</i> , 2009, 125, 217-229.	3.8	51
38	Associations among plasma metabolite levels and short-term exposure to PM2.5 and ozone in a cardiac catheterization cohort. <i>Environment International</i> , 2016, 97, 76-84.	10.0	51
39	Baseline Colonoscopy Findings Associated With 10-Year Outcomes in a Screening Cohort Undergoing Colonoscopy Surveillance. <i>Gastroenterology</i> , 2020, 158, 862-874.e8.	1.3	51
40	Ordered subset linkage analysis supports a susceptibility locus for age-related macular degeneration on chromosome 16p12. <i>BMC Genetics</i> , 2004, 5, 18.	2.7	48
41	The APL Test: Extension to General Nuclear Families and Haplotypes and Examination of Its Robustness. <i>Human Heredity</i> , 2006, 61, 189-199.	0.8	48
42	Extension of the SIMLA Package for Generating Pedigrees with Complex Inheritance Patterns: Environmental Covariates, Gene-Gene and Gene-Environment Interaction. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2005, 4, Article15.	0.6	47
43	Metabolic profiles predict adverse events after coronary artery bypass grafting. <i>Journal of Thoracic and Cardiovascular Surgery</i> , 2012, 143, 873-878.	0.8	45
44	Evaluating DNA methylation age on the Illumina MethylationEPIC Bead Chip. <i>PLoS ONE</i> , 2019, 14, e0207834.	2.5	44
45	Genomic convergence: identifying candidate genes for Parkinson's disease by combining serial analysis of gene expression and genetic linkage. <i>Human Molecular Genetics</i> , 2003, 12, 671-7.	2.9	44
46	A common variant in the CDKN2B gene on chromosome 9p21 protects against coronary artery disease in Americans of African ancestry. <i>Journal of Human Genetics</i> , 2011, 56, 224-229.	2.3	43
47	Gene by stress genome-wide interaction analysis and path analysis identify EBF1 as a cardiovascular and metabolic risk gene. <i>European Journal of Human Genetics</i> , 2015, 23, 854-862.	2.8	42
48	Simple method of measurement of orotic acid and orotidine in urine. <i>Biomedical Applications</i> , 1989, 493, 388-391.	1.7	40
49	An atlas connecting shared genetic architecture of human diseases and molecular phenotypes provides insight into COVID-19 susceptibility. <i>Genome Medicine</i> , 2021, 13, 83.	8.2	40
50	GxE Interactions between FOXO Genotypes and Tea Drinking Are Significantly Associated with Cognitive Disability at Advanced Ages in China. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2015, 70, 426-433.	3.6	34
51	Clinical utility of a Web-enabled risk-assessment and clinical decision support program. <i>Genetics in Medicine</i> , 2016, 18, 1020-1028.	2.4	34
52	Reclassification of cardiovascular risk using integrated clinical and molecular biosignatures: Design of and rationale for the Measurement to Understand the Reclassification of Disease of Cabarrus and Kannapolis (MURDOCK) Horizon 1 Cardiovascular Disease Study. <i>American Heart Journal</i> , 2010, 160, 371-379.e2.	2.7	33
53	Aging-related atherosclerosis is exacerbated by arterial expression of tumor necrosis factor receptor-1: evidence from mouse models and human association studies. <i>Human Molecular Genetics</i> , 2010, 19, 2754-2766.	2.9	32
54	Genetic and functional association of FAM5C with myocardial infarction. <i>BMC Medical Genetics</i> , 2008, 9, 33.	2.1	31

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55	Increased hypothalamic [3H]flunitrazepam binding in hypothalamic-pituitary-adrenal axis hyporesponsive Lewis rats. <i>Brain Research</i> , 1992, 569, 295-299.	2.2	29
56	The Finland–United States Investigation of Non–Insulin–Dependent Diabetes Mellitus Genetics (FUSION) Study. II. An Autosomal Genome Scan for Diabetes–Related Quantitative Trait Loci. <i>American Journal of Human Genetics</i> , 2000, 67, 1186-1200.	6.2	28
57	Validation Study of Genetic Associations with Coronary Artery Disease on Chromosome 3q13–21 and Potential Effect Modification by Smoking. <i>Annals of Human Genetics</i> , 2009, 73, 551-558.	0.8	27
58	Association of Roadway Proximity with Fasting Plasma Glucose and Metabolic Risk Factors for Cardiovascular Disease in a Cross-Sectional Study of Cardiac Catheterization Patients. <i>Environmental Health Perspectives</i> , 2015, 123, 1007-1014.	6.0	27
59	Short-term effects of fine particulate matter and ozone on the cardiac conduction system in patients undergoing cardiac catheterization. <i>Particle and Fibre Toxicology</i> , 2018, 15, 38.	6.2	26
60	Polymorphic variants in tenascin-C (TNC) are associated with atherosclerosis and coronary artery disease. <i>Human Genetics</i> , 2011, 129, 641-654.	3.8	25
61	Brain-derived neurotrophic factor rs6265 (Val66Met) polymorphism is associated with disease severity and incidence of cardiovascular events in a patient cohort. <i>American Heart Journal</i> , 2017, 190, 40-45.	2.7	25
62	Genetic Variants in the Bone Morphogenic Protein Gene Family Modify the Association between Residential Exposure to Traffic and Peripheral Arterial Disease. <i>PLoS ONE</i> , 2016, 11, e0152670.	2.5	23
63	ALOX5AP variants are associated with in-stent restenosis after percutaneous coronary intervention. <i>Atherosclerosis</i> , 2008, 201, 148-154.	0.8	22
64	The genomic medicine model: an integrated approach to implementation of family health history in primary care. <i>Personalized Medicine</i> , 2013, 10, 295-306.	1.5	22
65	Genetic Simulation Tools for Post–Genome Wide Association Studies of Complex Diseases. <i>Genetic Epidemiology</i> , 2015, 39, 11-19.	1.3	22
66	Genetic analysis for common complex disease. <i>American Heart Journal</i> , 2000, 140, S36-S44.	2.7	21
67	Fine mapping of a linkage peak with integration of lipid traits identifies novel coronary artery disease genes on chromosome 5. <i>BMC Genetics</i> , 2012, 13, 12.	2.7	21
68	A Functional Polymorphism in the 5HTR2C Gene Associated with Stress Responses Also Predicts Incident Cardiovascular Events. <i>PLoS ONE</i> , 2013, 8, e82781.	2.5	21
69	Impact of Genetic Testing and Family Health History Based Risk Counseling on Behavior Change and Cognitive Precursors for Type 2 Diabetes. <i>Journal of Genetic Counseling</i> , 2017, 26, 133-140.	1.6	21
70	A genome-wide trans-ethnic interaction study links the PIGR-FCAMR locus to coronary atherosclerosis via interactions between genetic variants and residential exposure to traffic. <i>PLoS ONE</i> , 2017, 12, e0173880.	2.5	21
71	Genome–wide linkage analysis of quantitative biomarker traits of osteoarthritis in a large, multigenerational extended family. <i>Arthritis and Rheumatism</i> , 2010, 62, 781-790.	6.7	20
72	Neighborhood Sociodemographic Effects on the Associations Between Long-term PM2.5 Exposure and Cardiovascular Outcomes and Diabetes Mellitus. <i>Environmental Epidemiology</i> , 2019, 3, e038.	3.0	20

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73	Epigenome-wide association study of kidney function identifies trans-ethnic and ethnic-specific loci. <i>Genome Medicine</i> , 2021, 13, 74.	8.2	20
74	Accelerated epigenetic age as a biomarker of cardiovascular sensitivity to traffic-related air pollution. <i>Aging</i> , 2020, 12, 24141-24155.	3.1	18
75	Simultaneous Consideration of Multiple Candidate Protein Biomarkers for Long-Term Risk for Cardiovascular Events. <i>Circulation: Cardiovascular Genetics</i> , 2015, 8, 168-177.	5.1	17
76	A novel approach for measuring residential socioeconomic factors associated with cardiovascular and metabolic health. <i>Journal of Exposure Science and Environmental Epidemiology</i> , 2017, 27, 281-289.	3.9	17
77	The Gulf War Era Cohort and Biorepository: A Longitudinal Research Resource of Veterans of the 1990-1991 Gulf War Era. <i>American Journal of Epidemiology</i> , 2018, 187, 2279-2291.	3.4	17
78	Epigenetic Profiling Identifies Novel Genes for Ascending Aortic Aneurysm Formation with Bicuspid Aortic Valves. <i>Heart Surgery Forum</i> , 2015, 18, 134.	0.5	17
79	Associations Between Residential Proximity to Traffic and Vascular Disease in a Cardiac Catheterization Cohort. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2018, 38, 275-282.	2.4	15
80	Association between short-term exposure to ambient fine particulate matter and myocardial injury in the CATHGEN cohort. <i>Environmental Pollution</i> , 2021, 275, 116663.	7.5	15
81	Interpretation of simultaneous linkage and family-based association tests in genome screens. <i>Genetic Epidemiology</i> , 2007, 31, 134-142.	1.3	14
82	Interaction Between the <i>FOXO1A-209</i> Genotype and Tea Drinking Is Significantly Associated with Reduced Mortality at Advanced Ages. <i>Rejuvenation Research</i> , 2016, 19, 195-203.	1.8	14
83	Recommendations for Improving Identification and Quantification in Non-Targeted, GC-MS-Based Metabolomic Profiling of Human Plasma. <i>Metabolites</i> , 2017, 7, 45.	2.9	14
84	Gulf War illness in the Gulf War Era Cohort and Biorepository: The Kansas and Centers for Disease Control definitions. <i>Life Sciences</i> , 2021, 278, 119454.	4.3	14
85	Genome-Wide Linkage Analysis of Cardiovascular Disease Biomarkers in a Large, Multigenerational Family. <i>PLoS ONE</i> , 2013, 8, e71779.	2.5	12
86	A multi-population phenome-wide association study of genetically-predicted height in the Million Veteran Program. <i>PLoS Genetics</i> , 2022, 18, e1010193.	3.5	12
87	Ordered-subset analysis (OSA) for family-based association mapping of complex traits. <i>Genetic Epidemiology</i> , 2008, 32, 627-637.	1.3	10
88	Gene-smoking interactions in multiple Rho-GTPase pathway genes in an early-onset coronary artery disease cohort. <i>Human Genetics</i> , 2013, 132, 1371-1382.	3.8	10
89	Phenotyping clinical disorders: lessons learned from pelvic organ prolapse. <i>American Journal of Obstetrics and Gynecology</i> , 2013, 208, 360-365.	1.3	10
90	Association of standard clinical and laboratory variables with red blood cell distribution width. <i>American Heart Journal</i> , 2016, 174, 22-28.	2.7	10

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91	Effects of covariates: A summary of Group 5 contributions. <i>Genetic Epidemiology</i> , 2003, 25, S43-S49.	1.3	9
92	Rheumatoid arthritis T cell and muscle oxidative metabolism associate with exercise-induced changes in cardiorespiratory fitness. <i>Scientific Reports</i> , 2022, 12, 7450.	3.3	9
93	Ordered subset analysis for case-control studies. <i>Genetic Epidemiology</i> , 2010, 34, 407-417.	1.3	8
94	Computing a Synthetic Chronic Psychosocial Stress Measurement in Multiple Datasets and its Application in the Replication of G × E Interactions of the <i>EBF1</i> Gene. <i>Genetic Epidemiology</i> , 2015, 39, 489-497.	1.3	8
95	Apolipoprotein L1 Genetic Variants Are Associated with Chronic Kidney Disease but Not with Cardiovascular Disease in a Population Referred for Cardiac Catheterization. <i>CardioRenal Medicine</i> , 2017, 7, 96-103.	1.9	8
96	Genetic Variation in Acid Ceramidase Predicts Non-completion of an Exercise Intervention. <i>Frontiers in Physiology</i> , 2018, 9, 781.	2.8	8
97	Drebrin attenuates atherosclerosis by limiting smooth muscle cell transdifferentiation. <i>Cardiovascular Research</i> , 2022, 118, 772-784.	3.8	8
98	Linkage analysis with gene-environment interaction: model illustration and performance of ordered subset analysis. <i>Genetic Epidemiology</i> , 2006, 30, 409-422.	1.3	7
99	A general integrative genomic feature transcription factor binding site prediction method applied to analysis of USF 1 binding in cardiovascular disease. <i>Human Genomics</i> , 2009, 3, 221.	2.9	7
100	Genetic Colorectal Cancer and Adenoma Risk Variants Are Associated with Increasing Cumulative Adenoma Counts. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2269-2276.	2.5	7
101	Genomics of Gulf War Illness in U.S. Veterans Who Served during the 1990-1991 Persian Gulf War: Methods and Rationale for Veterans Affairs Cooperative Study #2006. <i>Brain Sciences</i> , 2021, 11, 845.	2.3	7
102	APOL1 risk alleles among individuals with CKD in Northern Tanzania: A pilot study. <i>PLoS ONE</i> , 2017, 12, e0181811.	2.5	7
103	Associations between neighborhood socioeconomic cluster and hypertension, diabetes, myocardial infarction, and coronary artery disease within a cohort of cardiac catheterization patients. <i>American Heart Journal</i> , 2022, 243, 201-209.	2.7	7
104	Interpreting analyses of continuous covariates in affected sibling pair linkage studies. <i>Genetic Epidemiology</i> , 2007, 31, 541-552.	1.3	6
105	Multistage designs in the genomic era: Providing balance in complex disease studies. <i>Genetic Epidemiology</i> , 2007, 31, S118-S123.	1.3	6
106	The genetic basis for survivorship in coronary artery disease. <i>Frontiers in Genetics</i> , 2013, 4, 191.	2.3	6
107	Case-Only Survival Analysis Reveals Unique Effects of Genotype, Sex, and Coronary Disease Severity on Survivorship. <i>PLoS ONE</i> , 2016, 11, e0154856.	2.5	6
108	Systolic Blood Pressure and Socioeconomic Status in a large multi-study population. <i>SSM - Population Health</i> , 2019, 9, 100498.	2.7	6



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109	Life After the Screen: Making Sense of Many P-values. <i>Genetic Epidemiology</i> , 2001, 21, S546-51.	1.3	5
110	Searching for epistatic interactions in nuclear families using conditional linkage analysis. <i>BMC Genetics</i> , 2005, 6, S148.	2.7	5
111	Increased Efficiency of Case-Control Association Analysis by Using Allele-Sharing and Covariate Information. <i>Human Heredity</i> , 2008, 65, 154-165.	0.8	5
112	Short-term effects of air temperature on plasma metabolite concentrations in patients undergoing cardiac catheterization. <i>Environmental Research</i> , 2016, 151, 224-232.	7.5	5
113	Lack of Association of a Functional Polymorphism in the Serotonin Receptor Gene With Body Mass Index and Depressive Symptoms in a Large Meta-Analysis of Population Based Studies. <i>Frontiers in Genetics</i> , 2018, 9, 423.	2.3	5
114	Research tool for classifying Gulf War illness using survey responses: Lessons for writing replicable algorithms for symptom-based conditions. <i>Life Sciences</i> , 2021, 282, 119808.	4.3	5
115	Angiotensin-converting enzyme gene insertion/deletion polymorphism and cardiovascular disease: Identifying the guideposts for navigating the genetics landscape. <i>American Heart Journal</i> , 2002, 144, 747-749.	2.7	5
116	Gene-Toxicant Interactions in Gulf War Illness: Differential Effects of the PON1 Genotype. <i>Brain Sciences</i> , 2021, 11, 1558.	2.3	5
117	858 Clinical Risk Group at Baseline Is Associated With 10 Year Outcomes in a Screening Cohort-Longitudinal Analysis of the CSP 380 Cohort. <i>Gastroenterology</i> , 2016, 150, S184.	1.3	4
118	High-Risk Adenomas at Screening Colonoscopy Remain Predictive of Future High-Risk Adenomas Despite an Intervening Negative Colonoscopy. <i>American Journal of Gastroenterology</i> , 2020, 115, 1275-1282.	0.4	4
119	Genetic Variants Associated with Vein Graft Stenosis after Coronary Artery Bypass Grafting. <i>Heart Surgery Forum</i> , 2015, 18, 001.	0.5	4
120	Adjusting for covariates on a slippery slope: linkage analysis of change over time. <i>BMC Genetics</i> , 2003, 4, S50.	2.7	3
121	Visualizing genotype-phenotype relationships in the GAW15 simulated data. <i>BMC Proceedings</i> , 2007, 1, S132.	1.6	3
122	Comparison of GIST and LAMP on the GAW15 simulated data. <i>BMC Proceedings</i> , 2007, 1, S41.	1.6	3
123	Assessment of LD Matrix Measures for the Analysis of Biological Pathway Association. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2010, 9, Article35.	0.6	3
124	Developing a synthetic psychosocial stress measure and harmonizing CVD-risk data: a way forward to GxE meta- and mega-analyses. <i>BMC Research Notes</i> , 2018, 11, 504.	1.4	3
125	Th17 Immunity in the Colon Is Controlled by Two Novel Subsets of Colon-Specific Mononuclear Phagocytes. <i>Frontiers in Immunology</i> , 2021, 12, 661290.	4.8	3
126	Genome-Wide Variants Associated With Longitudinal Survival Outcomes Among Individuals With Coronary Artery Disease. <i>Frontiers in Genetics</i> , 2021, 12, 661497.	2.3	3



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127	Health-Related Quality of Life by Gulf War Illness Case Status. <i>International Journal of Environmental Research and Public Health</i> , 2022, 19, 4425.	2.6	3
128	Exposures to low-levels of fine particulate matter are associated with acute changes in heart rate variability, cardiac repolarization, and circulating blood lipids in coronary artery disease patients. <i>Environmental Research</i> , 2022, 214, 113768.	7.5	3
129	Two-stage study designs for analyzing disease-associated covariates: linkage thresholds and case-selection strategies. <i>BMC Proceedings</i> , 2007, 1, S138.	1.6	2
130	Cigarette smoking status has a modifying effect on the association between polymorphisms in KALRN and measures of cardiovascular risk in the diabetes heart study. <i>Genes and Genomics</i> , 2011, 33, 483-490.	1.4	2
131	Characterizing chronological accumulation of comorbidities in healthy veterans: a computational approach. <i>Scientific Reports</i> , 2021, 11, 8104.	3.3	2
132	Ten or More Cumulative Lifetime Adenomas Are Associated with Increased Risk for Advanced Neoplasia and Colorectal Cancer. <i>Digestive Diseases and Sciences</i> , 2022, 67, 2526-2534.	2.3	2
133	Screening Colonoscopy Findings Are Associated With Noncolorectal Cancer Mortality. <i>Clinical and Translational Gastroenterology</i> , 2022, 13, e00479.	2.5	2
134	Pedigree Selection and Information Content. <i>Current Protocols in Human Genetics</i> , 2001, 29, Unit 1.2.	3.5	1
135	Evaluating the precision of EBF1 SNP x stress interaction association: sex, race, and age differences in a big harmonized data set of 28,026 participants. <i>Translational Psychiatry</i> , 2020, 10, 351.	4.8	1
136	Sex-dimorphic gene effects on survival outcomes in people with coronary artery disease. <i>American Heart Journal Plus</i> , 2022, 17, 100152.	0.6	1
137	Nonparametric Linkage Analysis. , 0, , 283-328.		0
138	Mo1724 Risk Factors Associated With the Development of Adenoma Multiplicity in a Screening Cohort. <i>Gastroenterology</i> , 2016, 150, S763.	1.3	0
139	Colorectal Cancer Risk Factors in Veterans with and Without Adenoma Multiplicity in a Screening Cohort. <i>Gastroenterology</i> , 2017, 152, S543-S544.	1.3	0
140	Risk factors for interval advanced colorectal neoplasia after screening colonoscopy.. <i>Journal of Clinical Oncology</i> , 2015, 33, 3539-3539.	1.6	0
141	Validation of the NCI colorectal cancer risk assessment tool in the CSP 380 veterans cohort.. <i>Journal of Clinical Oncology</i> , 2017, 35, e15135-e15135.	1.6	0
142	Validation of the NCI Colorectal Cancer Risk Assessment Tool for baseline advanced neoplasia in a veterans cohort.. <i>Journal of Clinical Oncology</i> , 2019, 37, 521-521.	1.6	0
143	Characterization of temporal relationships of comorbidities developed following cancer diagnoses in veterans.. <i>Journal of Clinical Oncology</i> , 2019, 37, e18049-e18049.	1.6	0
144	Abstract 18660: CVSN Best Abstract Award: Genome-wide Candidates Unique to Females With Coronary Artery Disease Significantly Predict Mortality Risk. <i>Circulation</i> , 2015, 132, .	1.6	0