Ping Zeng

List of Publications by Citations

Source: https://exaly.com/author-pdf/8478738/ping-zeng-publications-by-citations.pdf

Version: 2024-04-28

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.

59	543	13	21
papers	citations	h-index	g-index
72	934	5.6	4.81
ext. papers	ext. citations	avg, IF	L-index

#	Paper	IF	Citations
59	Detecting epistasis with the marginal epistasis test in genetic mapping studies of quantitative traits. <i>PLoS Genetics</i> , 2017 , 13, e1006869	6	53
58	Non-parametric genetic prediction of complex traits with latent Dirichlet process regression models. <i>Nature Communications</i> , 2017 , 8, 456	17.4	53
57	Causal effects of blood lipids on amyotrophic lateral sclerosis: a Mendelian randomization study. <i>Human Molecular Genetics</i> , 2019 , 28, 688-697	5.6	52
56	Statistical analysis for genome-wide association study. <i>Journal of Biomedical Research</i> , 2015 , 29, 285-97	1.5	40
55	Causal association of type 2 diabetes with amyotrophic lateral sclerosis: new evidence from Mendelian randomization using GWAS summary statistics. <i>BMC Medicine</i> , 2019 , 17, 225	11.4	35
54	Testing and controlling for horizontal pleiotropy with probabilistic Mendelian randomization in transcriptome-wide association studies. <i>Nature Communications</i> , 2020 , 11, 3861	17.4	27
53	Causal Association Between Birth Weight and Adult Diseases: Evidence From a Mendelian Randomization Analysis. <i>Frontiers in Genetics</i> , 2019 , 10, 618	4.5	24
52	Identifying and exploiting trait-relevant tissues with multiple functional annotations in genome-wide association studies. <i>PLoS Genetics</i> , 2018 , 14, e1007186	6	23
51	Pleiotropic mapping and annotation selection in genome-wide association studies with penalized Gaussian mixture models. <i>Bioinformatics</i> , 2018 , 34, 2797-2807	7.2	17
50	Prediction of gene expression with cis-SNPs using mixed models and regularization methods. <i>BMC Genomics</i> , 2017 , 18, 368	4.5	17
49	Stray dogs as indicators of Toxoplasma gondii distributed in the environment: the first report across an urban-rural gradient in China. <i>Parasites and Vectors</i> , 2012 , 5, 5	4	17
48	Alcohol Drinking and Amyotrophic Lateral Sclerosis: An Instrumental Variable Causal Inference. <i>Annals of Neurology</i> , 2020 , 88, 195-198	9.4	16
47	Variable selection approach for zero-inflated count data via adaptive lasso. <i>Journal of Applied Statistics</i> , 2014 , 41, 879-894	1	16
46	Multi-trait transcriptome-wide association studies with probabilistic Mendelian randomization. <i>American Journal of Human Genetics</i> , 2021 , 108, 240-256	11	13
45	Association Between Premorbid Body Mass Index and Amyotrophic Lateral Sclerosis: Causal Inference Through Genetic Approaches. <i>Frontiers in Neurology</i> , 2019 , 10, 543	4.1	12
44	Likelihood ratio tests in rare variant detection for continuous phenotypes. <i>Annals of Human Genetics</i> , 2014 , 78, 320-32	2.2	12
43	Relationship between birth weight and chronic kidney disease: evidence from systematics review and two-sample Mendelian randomization analysis. <i>Human Molecular Genetics</i> , 2020 , 29, 2261-2274	5.6	10

(2021-2015)

42	Permutation-based variance component test in generalized linear mixed model with application to multilocus genetic association study. <i>BMC Medical Research Methodology</i> , 2015 , 15, 37	4.7	7
41	Rare variants detection with kernel machine learning based on likelihood ratio test. <i>PLoS ONE</i> , 2014 , 9, e93355	3.7	7
40	Evaluation of the causal effects of blood lipid levels on gout with summary level GWAS data: two-sample Mendelian randomization and mediation analysis. <i>Journal of Human Genetics</i> , 2021 , 66, 465	5-473	7
39	Statistical methods for mediation analysis in the era of high-throughput genomics: Current successes and future challenges. <i>Computational and Structural Biotechnology Journal</i> , 2021 , 19, 3209-32	224 ⁸	7
38	Jackknife Model Averaging Prediction Methods for Complex Phenotypes with Gene Expression Levels by Integrating External Pathway Information. <i>Computational and Mathematical Methods in Medicine</i> , 2019 , 2019, 2807470	2.8	6
37	Multiple-Tissue Integrative Transcriptome-Wide Association Studies Discovered New Genes Associated With Amyotrophic Lateral Sclerosis. <i>Frontiers in Genetics</i> , 2020 , 11, 587243	4.5	6
36	Birth Weight and Stroke in Adult Life: Genetic Correlation and Causal Inference With Genome-Wide Association Data Sets. <i>Frontiers in Neuroscience</i> , 2020 , 14, 479	5.1	5
35	A two-stage Bayesian method for estimating accuracy and disease prevalence for two dependent dichotomous screening tests when the status of individuals who are negative on both tests is unverified. <i>BMC Medical Research Methodology</i> , 2014 , 14, 110	4.7	5
34	How Can Gene-Expression Information Improve Prognostic Prediction in TCGA Cancers: An Empirical Comparison Study on Regularization and Mixed Cox Models. <i>Frontiers in Genetics</i> , 2020 , 11, 920	4.5	5
33	Aggregating multiple expression prediction models improves the power of transcriptome-wide association studies. <i>Human Molecular Genetics</i> , 2021 , 30, 939-951	5.6	5
32	Assessing the Relationship Between Leukocyte Telomere Length and Cancer Risk/Mortality in UK Biobank and TCGA Datasets With the Genetic Risk Score and Mendelian Randomization Approaches. <i>Frontiers in Genetics</i> , 2020 , 11, 583106	4.5	4
31	Birth weight is not causally associated with adult asthma: results from instrumental variable analyses. <i>Scientific Reports</i> , 2019 , 9, 7647	4.9	3
30	Verification of the correlation between progression-free survival and overall survival considering magnitudes of survival post- progression in the treatment of four types of cancer. <i>Asian Pacific Journal of Cancer Prevention</i> , 2015 , 16, 1001-6	1.7	3
29	Testing and controlling for horizontal pleiotropy with the probabilistic Mendelian randomization in transcriptome-wide association studies		3
28	Are blood lipids risk factors for fracture? Integrative evidence from instrumental variable causal inference and mediation analysis using genetic data. <i>Bone</i> , 2020 , 131, 115174	4.7	3
27	Detection of Genetic Overlap Between Rheumatoid Arthritis and Systemic Lupus Erythematosus Using GWAS Summary Statistics. <i>Frontiers in Genetics</i> , 2021 , 12, 656545	4.5	3
26	Birth weight is positively associated with adult osteoporosis risk: observational and Mendelian randomization studies. <i>Journal of Bone and Mineral Research</i> , 2021 , 36, 1469-1480	6.3	3
25	Evaluating marginal genetic correlation of associated loci for complex diseases and traits between European and East Asian populations. <i>Human Genetics</i> , 2021 , 140, 1285-1297	6.3	3

24	New novel non-MHC genes were identified for cervical cancer with an integrative analysis approach of transcriptome-wide association study. <i>Journal of Cancer</i> , 2021 , 12, 840-848	4.5	3
23	Detecting the Genomic Signature of Divergent Selection in Presence of Gene Flow. <i>Current Genomics</i> , 2015 , 16, 194-202	2.6	2
22	Leveraging methylation to identify the potential causal genes associated with survival in lung adenocarcinoma and lung squamous cell carcinoma. <i>Oncology Letters</i> , 2020 , 20, 193-200	2.6	2
21	Leveraging Methylation Alterations to Discover Potential Causal Genes Associated With the Survival Risk of Cervical Cancer in TCGA Through a Two-Stage Inference Approach. <i>Frontiers in Genetics</i> , 2021 , 12, 667877	4.5	2
20	Mendelian Randomization Highlights the Causal Role of Normal Thyroid Function on Blood Lipid Profiles. <i>Endocrinology</i> , 2021 , 162,	4.8	2
19	Evaluation of vaccine seroresponse rates and adverse event rates through Bayesian and frequentist methods. <i>Human Vaccines and Immunotherapeutics</i> , 2015 , 11, 1557-63	4.4	1
18	Cis-SNPs Set Testing and PrediXcan Analysis for Gene Expression Data using Linear Mixed Models. <i>Scientific Reports</i> , 2017 , 7, 15237	4.9	1
17	Stability SCAD: a powerful approach to detect interactions in large-scale genomic study. <i>BMC Bioinformatics</i> , 2014 , 15, 62	3.6	1
16	Causal Association between Birth Weight and Adult Diseases: Evidence from a Mendelian Randomisation Analysis		1
15	Detecting Epistasis with the Marginal Epistasis Test in Genetic Mapping Studies of Quantitative Traits		1
14	Mendelian randomization implies no direct causal association between leukocyte telomere length and amyotrophic lateral sclerosis. <i>Scientific Reports</i> , 2020 , 10, 12184	4.9	1
13	Instrumental Heterogeneity in Sex-Specific Two-Sample Mendelian Randomization: Empirical Results From the Relationship Between Anthropometric Traits and Breast/Prostate Cancer. <i>Frontiers in Genetics</i> , 2021 , 12, 651332	4.5	1
12	IUSMMT: Survival mediation analysis of gene expression with multiple DNA methylation exposures and its application to cancers of TCGA. <i>PLoS Computational Biology</i> , 2021 , 17, e1009250	5	1
11	Identifying pleiotropic genes for complex phenotypes with summary statistics from a perspective of composite null hypothesis testing. <i>Briefings in Bioinformatics</i> , 2021 ,	13.4	1
10	A comprehensive gene-centric pleiotropic association analysis for 14 psychiatric disorders with GWAS summary statistics <i>BMC Medicine</i> , 2021 , 19, 314	11.4	О
9	Integrative eQTL-weighted hierarchical Cox models for SNP-set based time-to-event association studies. <i>Journal of Translational Medicine</i> , 2021 , 19, 418	8.5	O
8	Evaluating Causal Relationship Between Metabolites and Six Cardiovascular Diseases Based on GWAS Summary Statistics. <i>Frontiers in Genetics</i> , 2021 , 12, 746677	4.5	0
7	Improved Detection of Potentially Pleiotropic Genes in Coronary Artery Disease and Chronic Kidney Disease Using GWAS Summary Statistics. <i>Frontiers in Genetics</i> , 2020 , 11, 592461	4.5	O

LIST OF PUBLICATIONS

6	Novel Autophagy-Related Gene Signature Investigation for Patients With Oral Squamous Cell Carcinoma. <i>Frontiers in Genetics</i> , 2021 , 12, 673319	4.5	O
5	Type 2 Diabetes Mellitus and Amyotrophic Lateral Sclerosis: Genetic Overlap, Causality, and Mediation. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021 , 106, e4497-e4508	5.6	0
4	Developing ZNF Gene Signatures Predicting Radiosensitivity of Patients with Breast Cancer. Journal of Oncology, 2021 , 2021, 9255494	4.5	0
3	Assessing Causal Relationship Between Human Blood Metabolites and Five Neurodegenerative Diseases With GWAS Summary Statistics <i>Frontiers in Neuroscience</i> , 2021 , 15, 680104	5.1	O
2	The Mediating Role of Social Support in the Relationship Between Psychological Capital and Depression Among Chinese Emergency Physicians <i>Psychology Research and Behavior Management</i> , 2022 , 15, 977-990	3.8	0
1	Exploring the association between birthweight and breast cancer using summary statistics from a perspective of genetic correlation, mediation, and causality <i>Journal of Translational Medicine</i> , 2022 , 20, 227	8.5	O