## Dabao Zhang

List of Publications by Year in descending order

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687363 434195 1,843 31 13 31 citations h-index g-index papers 31 31 31 2875 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Risk Analysis via Generalized Pareto Distributions. Journal of Business and Economic Statistics, 2022, 40, 852-867.	2.9	3
2	Serum metabolomic analysis reveals several novel metabolites in association with excessive alcohol use – an exploratory study. Translational Research, 2022, 240, 87-98.	5.0	3
3	Genetic architecture of root and shoot ionomes in rice (Oryza sativa L.). Theoretical and Applied Genetics, 2021, 134, 2613-2637.	3.6	9
4	Integrating Biological Knowledge Into Case–Control Analysis Through Iterated Conditional Modes/Medians Algorithm. Journal of Computational Biology, 2020, 27, 1171-1179.	1.6	2
5	Largeâ€scale identification of expression quantitative trait loci in Arabidopsis reveals novel candidate regulators of immune responses and other processes. Journal of Integrative Plant Biology, 2020, 62, 1469-1484.	8.5	7
6	Inferring Gene Regulatory Networks from a Population of Yeast Segregants. Scientific Reports, 2019, 9, 1197.	3.3	11
7	Advanced Statistical Methods for NMR-Based Metabolomics. Methods in Molecular Biology, 2019, 2037, 471-482.	0.9	1
8	A Coefficient of Determination for Generalized Linear Models. American Statistician, 2017, 71, 310-316.	1.6	226
9	Altered metabolite levels and correlations in patients with colorectal cancer and polyps detected using seemingly unrelated regression analysis. Metabolomics, 2017, 13, 1.	3.0	15
10	Exploring the Effects of Genetic Variants on Clinical Profiles of Parkinson's Disease Assessed by the Unified Parkinson's Disease Rating Scale and the Hoehn–Yahr Stage. PLoS ONE, 2016, 11, e0155758.	2.5	12
11	Generalized orthogonal components regression for high dimensional generalized linear models. Computational Statistics and Data Analysis, 2015, 88, 119-127.	1.2	1
12	Exploring Metabolic Profile Differences between Colorectal Polyp Patients and Controls Using Seemingly Unrelated Regression. Journal of Proteome Research, 2015, 14, 2492-2499.	3.7	28
13	Transcriptional activation of PRMT5 by NF-Y is required for cell growth and negatively regulated by the PKC/c-Fos signaling in prostate cancer cells. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2014, 1839, 1330-1340.	1.9	40
14	Lack of association between dendritic cell nuclear protein-1 gene and major depressive disorder in the Han Chinese population. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2013, 45, 7-10.	4.8	2
15	EMPIRICAL-LIKELIHOOD-BASED CONFIDENCE INTERVALS FOR CONDITIONAL VARIANCE IN HETEROSKEDASTIC REGRESSION MODELS. Econometric Theory, 2011, 27, 154-177.	0.7	11
16	Genome-wide case-control study in GAW17 using coalesced rare variants. BMC Proceedings, 2011, 5, S110.	1.6	4
17	Chromosome Size in Diploid Eukaryotic Species Centers on the Average Length with a Conserved Boundary. Molecular Biology and Evolution, 2011, 28, 1901-1911.	8.9	31
18	Genotyping Error Detection in Samples of Unrelated Individuals without Replicate Genotyping. Human Heredity, 2009, 67, 154-162.	0.8	7

#	Article	IF	CITATIONS
19	Penalized orthogonal-components regression for large p small n data. Electronic Journal of Statistics, 2009, 3, .	0.7	18
20	Nonparametric estimation of the dependence function for a multivariate extreme value distribution. Journal of Multivariate Analysis, 2008, 99, 577-588.	1.0	46
21	Variable selection for large p small n regression models with incomplete data: Mapping QTL with epistases. BMC Bioinformatics, 2008, 9, 251.	2.6	37
22	Proteomic analysis of peripheral leukocytes in Alzheimer's disease patients treated with divalproex sodium. Neurobiology of Aging, 2008, 29, 1631-1643.	3.1	23
23	Two-Dimensional Correlation Optimized Warping Algorithm for Aligning GC×GCâ^'MS Data. Analytical Chemistry, 2008, 80, 2664-2671.	6.5	93
24	Bayesian profiling of molecular signatures to predict event times. Theoretical Biology and Medical Modelling, 2007, 4, 3.	2.1	4
25	Multiplicative background correction for spotted microarrays to improve reproducibility. Genetical Research, 2006, 87, 195-206.	0.9	11
26	Orphanin FQ Antagonizes the Inhibition of Ca <sup>2+</sup> Currents Induced by Âμ-Opioid Receptors. Journal of Molecular Neuroscience, 2005, 25, 021-028.	2.3	12
27	Hierarchical Graphical Models. Journal of the American Statistical Association, 2005, 100, 719-727.	3.1	3
28	Mapping Multiple Quantitative Trait Loci by Bayesian Classification. Genetics, 2005, 169, 2305-2318.	2.9	41
29	HIV-1 Transactivator of Transcription Protein Induces Mitochondrial Hyperpolarization and Synaptic Stress Leading to Apoptosis. Journal of Immunology, 2005, 174, 4333-4344.	0.8	95
30	Bayesian Normalization and Identification for Differential Gene Expression Data. Journal of Computational Biology, 2005, 12, 391-406.	1.6	13
31	The Collaborative Cross, a community resource for the genetic analysis of complex traits. Nature Genetics, 2004, 36, 1133-1137.	21.4	1,034