Zhongxue Chen

List of Publications by Year in descending order

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Version: 2024-02-01

108 papers 2,633 citations

30 h-index 223531 46 g-index

109 all docs

109 docs citations

109 times ranked 3567 citing authors

#	Article	IF	CITATIONS
1	Social capital and disaster preparedness among low income Mexican Americans in a disaster prone area. Social Science and Medicine, 2013, 83, 50-60.	1.8	136
2	Feature mining and pattern classification for steganalysis of LSB matching steganography in grayscale images. Pattern Recognition, 2008, 41, 56-66.	5.1	106
3	An outcome analysis of endovascular versus open repair of blunt traumatic aortic injuries. Journal of Vascular Surgery, 2013, 57, 108-115.	0.6	99
4	The Argatroban and Tissue-Type Plasminogen Activator Stroke Study. Stroke, 2012, 43, 770-775.	1.0	93
5	A distribution free summarization method for Affymetrix GeneChip(R) arrays. Bioinformatics, 2007, 23, 321-327.	1.8	89
6	Gene selection and classification for cancer microarray data based on machine learning and similarity measures. BMC Genomics, 2011, 12, S1.	1.2	81
7	Predictors of youth e-cigarette use susceptibility in a U.S. nationally representative sample. Addictive Behaviors, 2018, 82, 79-85.	1.7	77
8	Insulin-degrading enzyme is exported via an unconventional protein secretion pathway. Molecular Neurodegeneration, 2009, 4, 4.	4.4	76
9	Image complexity and feature mining for steganalysis of least significant bit matching steganography. Information Sciences, 2008, 178, 21-36.	4.0	74
10	Independent component analysis of Alzheimer's DNA microarray gene expression data. Molecular Neurodegeneration, 2009, 4, 5.	4.4	72
11	Seafood Consumption and Blood Mercury Concentrations in Jamaican Children With and Without Autism Spectrum Disorders. Neurotoxicity Research, 2013, 23, 22-38.	1.3	59
12	Is the weighted z-test the best method for combining probabilities from independent tests?. Journal of Evolutionary Biology, 2011, 24, 926-930.	0.8	55
13	An improved approach to steganalysis of JPEG images. Information Sciences, 2010, 180, 1643-1655.	4.0	54
14	The role of drinking water sources, consumption of vegetables and seafood in relation to blood arsenic concentrations of Jamaican children with and without Autism Spectrum Disorders. Science of the Total Environment, 2012, 433, 362-370.	3.9	54
15	Association of Acute Increase in Plasma Neurofilament Light with Repetitive Subconcussive Head Impacts: A Pilot Randomized Control Trial. Journal of Neurotrauma, 2019, 36, 548-553.	1.7	51
16	Statistical methods on detecting differentially expressed genes for RNA-seq data. BMC Systems Biology, 2011, 5, S1.	3.0	49
17	A Longitudinal Study to Identify Laboratory Predictors of Liver Disease Outcome in Alagille Syndrome. Journal of Pediatric Gastroenterology and Nutrition, 2010, 50, 526-530.	0.9	47
18	A new statistical approach to combining p-values using gamma distribution and its application to genome-wide association study. BMC Bioinformatics, 2014, 15, S3.	1.2	46

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19	Feature Selection and Classification of MAQC-II Breast Cancer and Multiple Myeloma Microarray Gene Expression Data. PLoS ONE, 2009, 4, e8250.	1.1	45
20	Serial Transcriptome Analysis and Cross-Species Integration Identifies <i>Centromere-Associated Protein E</i> as a Novel Neuroblastoma Target. Cancer Research, 2010, 70, 2749-2758.	0.4	45
21	Endovascular treatment of hemodialysis access pseudoaneurysms. Journal of Vascular Surgery, 2012, 55, 1058-1062.	0.6	45
22	Identification of genes and pathways involved in kidney renal clear cell carcinoma. BMC Bioinformatics, 2014, 15, S2.	1.2	45
23	Safety and Tolerability of Lactobacillus reuteri DSM 17938 and Effects on Biomarkers in Healthy Adults: Results from a Randomized Masked Trial. PLoS ONE, 2012, 7, e43910.	1.1	44
24	The emerging genomics and systems biology research lead to systems genomics studies. BMC Genomics, 2014, 15, 11.	1.2	39
25	Comparison of feature selection and classification for MALDI-MS data. BMC Genomics, 2009, 10, S3.	1.2	37
26	Improved Approaches with Calibrated Neighboring Joint Density to Steganalysis and Seam-Carved Forgery Detection in JPEG Images. ACM Transactions on Intelligent Systems and Technology, 2015, 5, 1-30.	2.9	35
27	Maternal and Paternal Age are Jointly Associated with Childhood Autism in Jamaica. Journal of Autism and Developmental Disorders, 2012, 42, 1928-1938.	1.7	34
28	Perioperative assessment of platelet function by Thromboelastograph \hat{A}^{\otimes} Platelet Mapping \hat{a} , \hat{c} in cardiovascular patients undergoing non-cardiac surgery. Journal of Thrombosis and Thrombolysis, 2013, 35, 23-30.	1.0	34
29	Subcomplex Iλ Specifically Controls Integrated Mitochondrial Functions in Caenorhabditis elegans. PLoS ONE, 2009, 4, e6607.	1.1	32
30	Parameter Estimation for the Exponential-Normal Convolution Model for Background Correction of Affymetrix GeneChip Data. Statistical Applications in Genetics and Molecular Biology, 2006, 5, Article24.	0.2	31
31	Intention to Comply With Mandatory Hurricane Evacuation Orders Among Persons Living Along a Coastal Area. Disaster Medicine and Public Health Preparedness, 2013, 7, 46-54.	0.7	31
32	On the optimally weighted <mml:math altimg="si137.gif" display="inline" overflow="scroll" xmlns:mml="http://www.w3.org/1998/Math/MathML"><mml:mi>z</mml:mi></mml:math> -test for combining probabilities from independent studies. Computational Statistics and Data Analysis, 2014, 70, 387-394.	0.7	31
33	A Robust Method for Testing Association in Genome-Wide Association Studies. Human Heredity, 2012, 73, 26-34.	0.4	30
34	Neuro-Ophthalmologic Response to Repetitive Subconcussive Head Impacts. JAMA Ophthalmology, 2020, 138, 350.	1.4	30
35	Gender Differences in Tuberculosis Notification in Pakistan. American Journal of Tropical Medicine and Hygiene, 2011, 85, 514-517.	0.6	29
36	Detecting associated single-nucleotide polymorphisms on the X chromosome in case control genome-wide association studies. Statistical Methods in Medical Research, 2017, 26, 567-582.	0.7	28

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37	The association of depression stigma with barriers to seeking mental health care: a cross-sectional analysis. Journal of Mental Health, 2020, 29, 182-190.	1.0	28
38	A new statistical approach to detecting differentially methylated loci for case control Illumina array methylation data. Bioinformatics, 2012, 28, 1109-1113.	1.8	27
39	A new association test based on Chi-square partition for case-control GWA studies. Genetic Epidemiology, 2011, 35, 658-663.	0.6	22
40	The role of the folate pathway in pancreatic cancer risk. PLoS ONE, 2018, 13, e0193298.	1.1	22
41	A New Approach to Account for the Correlations among Single Nucleotide Polymorphisms in Genome-Wide Association Studies. Human Heredity, 2011, 72, 1-9.	0.4	20
42	Return to Learn: Academic Effects of Concussion in High School and College Student-Athletes. Frontiers in Pediatrics, 2020, 8, 57.	0.9	20
43	Clinicopathological risk factors for gastric cancer: a retrospective cohort study in China. BMJ Open, 2019, 9, e030639.	0.8	19
44	Age-adjusted nonparametric detection of differential DNA methylation with case-control designs. BMC Bioinformatics, 2013, 14, 86.	1.2	18
45	Association tests through combining -values for case control genome-wide association studies. Statistics and Probability Letters, 2013, 83, 1854-1862.	0.4	17
46	Goodness-of-fit test for meta-analysis. Scientific Reports, 2015, 5, 16983.	1.6	17
47	Testing for association in case-control genome-wide association studies with shared controls. Statistical Methods in Medical Research, 2016, 25, 954-967.	0.7	17
48	Design and analysis of multiple diseases genome-wide association studies without controls. Gene, 2012, 510, 87-92.	1.0	16
49	Perceptions of Social Support, Empowerment and Youth Risk Behaviors. Journal of Primary Prevention, 2012, 33, 33-46.	0.8	16
50	Pilot Study Comparing Total Intravenous Anesthesia to Inhalational Anesthesia in Endoscopic Sinus Surgery: Novel Approach of Blood Flow Quantification. Annals of Otology, Rhinology and Laryngology, 2012, 121, 725-732.	0.6	15
51	Misdiagnosis of Cerebellar Infarctions. Canadian Journal of Neurological Sciences, 2014, 41, 568-571.	0.3	15
52	Bayesian composite quantile regression. Journal of Statistical Computation and Simulation, 2015, 85, 3744-3754.	0.7	15
53	Comparison of Multiple Hazard Rate Functions. Biometrics, 2016, 72, 39-45.	0.8	15
54	Comparative study on serum levels of macro and trace elements in schizophrenia based on supervised learning methods. Journal of Trace Elements in Medicine and Biology, 2017, 43, 202-208.	1.5	15

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55	A Powerful Variant-Set Association Test Based on Chi-Square Distribution. Genetics, 2017, 207, 903-910.	1,2	15
56	Detection of JPEG double compression and identification of smartphone image source and post-capture manipulation. Applied Intelligence, 2013, 39, 705-726.	3.3	14
57	Preliminary evidence-based recommendations for return to learn: a novel pilot study tracking concussed college students. Concussion, 2019, 4, CNC63.	1.2	14
58	A distribution-free convolution model for background correction of oligonucleotide microarray data. BMC Genomics, 2009, 10, S19.	1.2	13
59	Identification of Smartphone-Image Source and Manipulation. Lecture Notes in Computer Science, 2012, , 262-271.	1.0	13
60	Detecting differentially methylated loci for Illumina Array methylation data based on human ovarian cancer data. BMC Medical Genomics, 2013, 6, S9.	0.7	13
61	Comparing survival curves based on medians. BMC Medical Research Methodology, 2016, 16, 33.	1.4	13
62	Geneâ€based sequential burden association test. Statistics in Medicine, 2019, 38, 2353-2363.	0.8	13
63	An improved robust association test for GWAS with multiple diseases. Statistics and Probability Letters, 2014, 91, 153-161.	0.4	12
64	Does idiopathic restless legs syndrome delay onset and reduce severity of Parkinson's disease: a pilot study. International Journal of Neuroscience, 2015, 125, 526-530.	0.8	12
65	Supervised learning-based tagSNP selection for genome-wide disease classifications. BMC Genomics, 2008, 9, S6.	1.2	11
66	Association Between Proteomic Blood Biomarkers and DTI/NODDI Metrics in Adolescent Football Players: A Pilot Study. Frontiers in Neurology, 2020, 11, 581781.	1,1	11
67	Genetic association test based on principal component analysis. Statistical Applications in Genetics and Molecular Biology, 2017, 16, 189-198.	0.2	10
68	Geneâ€based genetic association test with adaptive optimal weights. Genetic Epidemiology, 2018, 42, 95-103.	0.6	10
69	Extension of Mood's median test for survival data. Statistics and Probability Letters, 2014, 95, 77-84.	0.4	9
70	A new association test based on disease allele selection for case–control genome-wide association studies. BMC Genomics, 2014, 15, 358.	1,2	9
71	An improved two-stage procedure to compare hazard curves. Journal of Statistical Computation and Simulation, 2017, 87, 1877-1886.	0.7	9
72	Serum levels of chemical elements in esophageal squamous cell carcinoma in Anyang, China: a case-control study based on machine learning methods. BMJ Open, 2017, 7, e015443.	0.8	9

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73	A nonparametric test for equality of survival medians. Statistics in Medicine, 2012, 31, 844-854.	0.8	8
74	A note on Cochran test for homogeneity in one-way ANOVA and meta-analysis. Statistical Papers, 2014, 55, 301-310.	0.7	8
75	A gene-based test of association through an orthogonal decomposition of genotype scores. Human Genetics, 2017, 136, 1385-1394.	1.8	8
76	Comments on â€~Choosing an optimal method to combine <i>p</i> à€values' by Sungho Won, Nathan Morris, Qing Lu and Robert C. Elston, <i>Statistics in Medicine</i> 2009; 28 :1537–1553. Statistics in Medicine, 2011, 30, 2959-2961.	0.8	7
77	Advances in translational bioinformatics facilitate revealing the landscape of complex disease mechanisms. BMC Bioinformatics, 2014, 15, I1.	1.2	7
78	Brief Report of a Low-Cost Street-Corner Methodology Used to Assess Inner-City Residents' Awareness and Knowledge about Asthma. Journal of Urban Health, 2011, 88, 156-163.	1.8	6
79	A gene selection method for GeneChip array data with small sample sizes. BMC Genomics, 2011, 12, S7.	1.2	6
80	Assessment of gene order computing methods for Alzheimer's disease. BMC Medical Genomics, 2013, 6, S8.	0.7	6
81	A genetic association test through combining two independent tests. Genomics, 2019, 111, 1152-1159.	1.3	6
82	Testing for gene-gene interaction in case-control GWAS. Statistics and Its Interface, 2017, 10, 267-277.	0.2	6
83	Robust tests for combining p-values under arbitrary dependency structures. Scientific Reports, 2022, 12, 3158.	1.6	6
84	Identifying Differentially Expressed Genes based on probe level data for GeneChip arrays. International Journal of Computational Biology and Drug Design, 2010, 3, 237.	0.3	5
85	Detecting differentially methylated loci for multiple treatments based on high-throughput methylation data. BMC Bioinformatics, 2014, 15, 142.	1.2	5
86	CTP infarct core may predict poor outcome in stroke patients treated with IV t-PA. Journal of the Neurological Sciences, 2014, 340, 165-169.	0.3	5
87	A JPEG-based statistically invisible steganography. , 2011, , .		4
88	Inferences on correlation coefficients of bivariate log-normal distributions. Journal of Applied Statistics, 2015, 42, 603-613.	0.6	4
89	A novel gene-set association test based on variance-gamma distribution. Statistical Methods in Medical Research, 2019, 28, 2868-2875.	0.7	4
90	Acute Time-Course Changes in CCL11, CCL2, and IL-10 Levels After Controlled Subconcussive Head Impacts: A Pilot Randomized Clinical Trial. Journal of Head Trauma Rehabilitation, 2020, 35, 308-316.	1.0	4

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91	Optimal Tests for Combining p-Values. Applied Sciences (Switzerland), 2022, 12, 322.	1.3	4
92	Comments on †Application of an adaptive design to a randomized phase II selection trial in gastric cancer: a report of the study design†by Satoshi Morita and Junichi Sakamoto. <i>Pharmaceutical Statistics ⟨i⟩. Pharmaceutical Statistics, 2012, 11, 267-268.</i>	0.7	3
93	A nonparametric approach to detecting the difference of survival medians. Communications in Statistics Part B: Simulation and Computation, 2017, 46, 395-403.	0.6	3
94	Intention to Comply With Mandatory Hurricane Evacuation Orders Among Persons Living Along a Coastal Areaâ€"ERRATUM. Disaster Medicine and Public Health Preparedness, 2013, 7, 111-111.	0.7	2
95	Detecting methylation quantitative trait loci using a methylation random field method. Briefings in Bioinformatics, 2021, 22, .	3.2	2
96	CMAX3: A Robust Statistical Test for Genetic Association Accounting for Covariates. Genes, 2021, 12, 1723.	1.0	2
97	Microarray Gene Expression Classification Based on Supervised Learning and Similarity Measures. , 2006, , .		1
98	Feature Mining and Pattern Recognition in Multimedia Forensicsâ€"Detection of JPEG Image Based Steganography, Double-Compression, Interpolation and WAV Audio Based Steganography. , 2011, , 561-604.		1
99	No Association Observed between Coffee Intake and Risk of Non-Hodgkin Lymphoma among Postmenopausal Women. Journal of the Academy of Nutrition and Dietetics, 2022, 122, 1725-1736.	0.4	1
100	Media Reporting Bias Affects Reported Sudden Death Rates. Journal of the American College of Cardiology, 2011, 58, 990-991.	1.2	0
101	SS16. Endovascular Treatment of Hemodialysis Access Pseudoaneurysms: A Single Institution Experience. Journal of Vascular Surgery, 2011, 53, 24S.	0.6	0
102	Model for End-Stage Liver Disease Score is Associated With increased Transfusion Requirements in Patients Undergoing Orthotopic Liver Transplant. American Journal of Clinical Pathology, 2012, 138, A132-A132.	0.4	0
103	Improved survival after hemostatic resuscitation. Journal of Trauma, 2012, 72, 313.	2.3	0
104	Exposing Image Tampering with the Same Quantization Matrix., 2015,, 327-343.		0
105	Comparison of two methods in estimating standard error of the method of simulated moments estimators for generalized linear mixed models. Communications in Statistics Part B: Simulation and Computation, 2018, 47, 2886-2895.	0.6	0
106	Preliminary Evidence-Based Recommendations for Return to Learn: a novel pilot study tracking concussion recovery in college students. Neurology, 2019, 93, .	1.5	0
107	A Method to Detect AAC Audio Forgery. , 2015, , .		0
108	Association of tea-drinking habits with the risk of non-Hodgkin lymphoma: a prospective cohort study among postmenopausal women. British Journal of Nutrition, 2023, 129, 1543-1551.	1.2	0