

Yong Chen

List of Publications by Year in descending order

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Version: 2024-02-01

134
papers

2,326
citations

331538

21
h-index

302012

39
g-index

145
all docs

145
docs citations

145
times ranked

3180
citing authors

#	ARTICLE	IF	CITATIONS
1	Accounting for post-randomization variables in meta-analysis: A joint meta-regression approach. <i>Biometrics</i> , 2023, 79, 358-367.	0.8	0
2	Methodological challenges in spatial and contextual exposome-health studies. <i>Critical Reviews in Environmental Science and Technology</i> , 2023, 53, 827-846.	6.6	7
3	Heterogeneity-aware and communication-efficient distributed statistical inference. <i>Biometrika</i> , 2022, 109, 67-83.	1.3	24
4	EMBRACE: An EM-based bias reduction approach through Copas-model estimation for quantifying the evidence of selective publishing in network meta-analysis. <i>Biometrics</i> , 2022, 78, 754-765.	0.8	3
5	Controversy and Debate: Questionable utility of the relative risk in clinical research: Paper 2: Is the Odds Ratio "portable" in meta-analysis? Time to consider bivariate generalized linear mixed model. <i>Journal of Clinical Epidemiology</i> , 2022, 142, 280-287.	2.4	10
6	Controversy and Debate : Questionable utility of the relative risk in clinical research: Paper 4 :Odds Ratios are far from "portable" A call to use realistic models for effect variation in meta-analysis. <i>Journal of Clinical Epidemiology</i> , 2022, 142, 294-304.	2.4	10
7	Risk Factors for Diagnosis of Psoriatic Arthritis, Psoriasis, Rheumatoid Arthritis, and Ankylosing Spondylitis: A Set of Parallel Case-control Studies. <i>Journal of Rheumatology</i> , 2022, 49, 53-59.	1.0	19
8	Extending Hui-Walter framework to correlated outcomes with application to diagnosis tests of an eye disease among premature infants. <i>Statistics in Medicine</i> , 2022, 41, 433-448.	0.8	0
9	Privacy-preserving harmonization via distributed ComBat. <i>NeuroImage</i> , 2022, 248, 118822.	2.1	11
10	A Bayesian Network to Predict the Risk of Post Influenza Vaccination Guillain-Barré Syndrome: Development and Validation Study. <i>JMIR Public Health and Surveillance</i> , 2022, 8, e25658.	1.2	0
11	Does biologic therapy impact the development of PsA among patients with psoriasis?. <i>Annals of the Rheumatic Diseases</i> , 2022, 81, 80-86.	0.5	29
12	Analysis of Dual Combination Therapies Used in Treatment of Hypertension in a Multinational Cohort. <i>JAMA Network Open</i> , 2022, 5, e223877.	2.8	9
13	DLMM as a lossless one-shot algorithm for collaborative multi-site distributed linear mixed models. <i>Nature Communications</i> , 2022, 13, 1678.	5.8	9
14	SAT: a Surrogate-Assisted Two-wave case boosting sampling method, with application to EHR-based association studies. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2022, 29, 918-927.	2.2	2
15	ODACH: a one-shot distributed algorithm for Cox model with heterogeneous multi-center data. <i>Scientific Reports</i> , 2022, 12, 6627.	1.6	9
16	dPQL: a lossless distributed algorithm for generalized linear mixed model with application to privacy-preserving hospital profiling. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2022, 29, 1366-1371.	2.2	10
17	Accounting for publication bias using a bivariate trim and fill meta-analysis procedure. <i>Statistics in Medicine</i> , 2022, 41, 3466-3478.	0.8	8
18	Distributed Quasi-Poisson regression algorithm for modeling multi-site count outcomes in distributed data networks. <i>Journal of Biomedical Informatics</i> , 2022, 131, 104097.	2.5	9

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19	Federated Multi-view Learning for Private Medical Data Integration and Analysis. <i>ACM Transactions on Intelligent Systems and Technology</i> , 2022, 13, 1-23.	2.9	9
20	Distributed learning for heterogeneous clinical data with application to integrating COVID-19 data across 230 sites. <i>Npj Digital Medicine</i> , 2022, 5, .	5.7	8
21	The association of prescription opioid use with suicide attempts: An analysis of statewide medical claims data. <i>PLoS ONE</i> , 2022, 17, e0269809.	1.1	5
22	Multisite learning of high-dimensional heterogeneous data with applications to opioid use disorder study of 15,000 patients across 5 clinical sites. <i>Scientific Reports</i> , 2022, 12, .	1.6	0
23	Predictors of postinfectious inflammatory response syndrome in HIV-negative immunocompetent cryptococcal meningitis. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2021, 92, 680-681.	0.9	9
24	Serum antinuclear antibodies associate with worse prognosis in AQP4â€­positive neuromyelitis optica spectrum disorder. <i>Brain and Behavior</i> , 2021, 11, e01865.	1.0	12
25	Lossless integration of multiple electronic health records for identifying pleiotropy using summary statistics. <i>Nature Communications</i> , 2021, 12, 168.	5.8	2
26	A cost-effective chart review sampling design to account for phenotyping error in electronic health records (EHR) data. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2021, 29, 52-61.	2.2	5
27	Extracting postmarketing adverse events from safety reports in the vaccine adverse event reporting system (VAERS) using deep learning. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2021, 28, 1393-1400.	2.2	22
28	Prediction of postâ€­vaccination Guillainâ€­BarrÃ© syndrome using data from a passive surveillance system. <i>Pharmacoepidemiology and Drug Safety</i> , 2021, 30, 602-609.	0.9	7
29	Monitoring vaccine safety by studying temporal variation of adverse events using vaccine adverse event reporting system. <i>Annals of Applied Statistics</i> , 2021, 15, .	0.5	3
30	Estimating the reference interval from a fixed effects <sc>metaâ€­analysis</sc>. <i>Research Synthesis Methods</i> , 2021, 12, 630-640.	4.2	3
31	Variation in US Hospital Mortality Rates for Patients Admitted With COVID-19 During the First 6 Months of the Pandemic. <i>JAMA Internal Medicine</i> , 2021, 181, 471.	2.6	197
32	Using logic regression to characterize extreme heat exposures and their health associations: a time-series study of emergency department visits in Atlanta. <i>BMC Medical Research Methodology</i> , 2021, 21, 87.	1.4	5
33	Why Is the Electronic Health Record So Challenging for Research and Clinical Care?. <i>Methods of Information in Medicine</i> , 2021, 60, 032-048.	0.7	13
34	Studying pediatric health outcomes with electronic health records using Bayesian clustering and trajectory analysis. <i>Journal of Biomedical Informatics</i> , 2021, 113, 103654.	2.5	8
35	Predictive P-score for treatment ranking in Bayesian network meta-analysis. <i>BMC Medical Research Methodology</i> , 2021, 21, 213.	1.4	15
36	An efficient and accurate distributed learning algorithm for modeling multi-site zero-inflated count outcomes. <i>Scientific Reports</i> , 2021, 11, 19647.	1.6	9

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37	Global identifiability of latent class models with applications to diagnostic test accuracy studies: A Granger basis approach. <i>Biometrics</i> , 2020, 76, 98-108.	0.8	5
38	Embracing study heterogeneity for finding genetic interactions in large-scale research consortia. <i>Genetic Epidemiology</i> , 2020, 44, 52-66.	0.6	4
39	An augmented estimation procedure for EHR-based association studies accounting for differential misclassification. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2020, 27, 244-253.	2.2	9
40	Learning from electronic health records across multiple sites: A communication-efficient and privacy-preserving distributed algorithm. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2020, 27, 376-385.	2.2	61
41	A fast score test for generalized mixture models. <i>Biometrics</i> , 2020, 76, 811-820.	0.8	4
42	On specification tests for composite likelihood inference. <i>Biometrika</i> , 2020, 107, 907-917.	1.3	1
43	SCOR: A secure international informatics infrastructure to investigate COVID-19. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2020, 27, 1721-1726.	2.2	31
44	Testing small study effects in multivariate meta-analysis. <i>Biometrics</i> , 2020, 76, 1240-1250.	0.8	24
45	Learning from local to global: An efficient distributed algorithm for modeling time-to-event data. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2020, 27, 1028-1036.	2.2	46
46	Diagnostic accuracy of the PLASMIC score in patients with suspected thrombotic thrombocytopenic purpura: A systematic review and meta-analysis. <i>Transfusion</i> , 2020, 60, 2047-2057.	0.8	37
47	Risk of Persistent Opioid Use following Major Surgery in Matched Samples of Patients with and without Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2126-2133.	1.1	7
48	Impact of Hospitalization and Medication Switching on Post-discharge Adherence to Oral Anticoagulants in Patients With Atrial Fibrillation. <i>Pharmacotherapy</i> , 2020, 40, 1022-1035.	1.2	6
49	A historical review of publication bias. <i>Research Synthesis Methods</i> , 2020, 11, 725-742.	4.2	69
50	Ideas for how informaticians can get involved with COVID-19 research. <i>BioData Mining</i> , 2020, 13, 3.	2.2	20
51	A pseudolikelihood approach for assessing genetic association in case-control studies with unmeasured population structure. <i>Statistical Methods in Medical Research</i> , 2020, 29, 3153-3165.	0.7	0
52	Reducing Bias Due to Outcome Misclassification for Epidemiologic Studies Using EHR-derived Probabilistic Phenotypes. <i>Epidemiology</i> , 2020, Publish Ahead of Print, 542-550.	1.2	9
53	Back Cover Image. <i>Genetic Epidemiology</i> , 2020, 44, ii.	0.6	0
54	Electronic health records and polygenic risk scores for predicting disease risk. <i>Nature Reviews Genetics</i> , 2020, 21, 493-502.	7.7	78

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55	Semiparametric modelling and estimation of covariate-adjusted dependence between bivariate recurrent events. <i>Biometrics</i> , 2020, 76, 1229-1239.	0.8	2
56	How Computational Experiments Can Improve Our Understanding of the Genetic Architecture of Common Human Diseases. <i>Artificial Life</i> , 2020, 26, 23-37.	1.0	4
57	The Galaxy Plot: A New Visualization Tool for Bivariate Meta-Analysis Studies. <i>American Journal of Epidemiology</i> , 2020, 189, 861-869.	1.6	8
58	Use of Deep Learning to Analyze Social Media Discussions About the Human Papillomavirus Vaccine. <i>JAMA Network Open</i> , 2020, 3, e2022025.	2.8	32
59	Small-study effects: current practice and challenges for future research. <i>Statistics and Its Interface</i> , 2020, 13, 475-484.	0.2	5
60	Evaluation of Phenotyping Errors on Polygenic Risk Score Predictions. , 2020, , .		1
61	Does hospitalization for thromboembolism improve oral anticoagulant adherence in patients with atrial fibrillation?. <i>Journal of the American Pharmacists Association: JAPhA</i> , 2020, 60, 986-992.e2.	0.7	1
62	Robust-ODAL: Learning from heterogeneous health systems without sharing patient-level data. <i>Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing</i> , 2020, 25, 695-706.	0.7	7
63	Identifying Clinical Risk Factors for Opioid Use Disorder using a Distributed Algorithm to Combine Real-World Data from a Large Clinical Data Research Network. <i>AMIA ... Annual Symposium proceedings</i> , 2020, 2020, 1220-1229.	0.2	1
64	Leverage Real-world Longitudinal Data in Large Clinical Research Networks for Alzheimer's Disease and Related Dementia (ADRD). <i>AMIA ... Annual Symposium proceedings</i> , 2020, 2020, 393-401.	0.2	1
65	Copas-like selection model to correct publication bias in systematic review of diagnostic test studies. <i>Statistical Methods in Medical Research</i> , 2019, 28, 2912-2923.	0.7	5
66	Comparing drug safety of hepatitis C therapies using post-market data. <i>BMC Medical Informatics and Decision Making</i> , 2019, 19, 147.	1.5	7
67	Prediction of 30-day pediatric unplanned hospitalizations using the Johns Hopkins Adjusted Clinical Groups risk adjustment system. <i>PLoS ONE</i> , 2019, 14, e0221233.	1.1	15
68	A regression framework to uncover pleiotropy in large-scale electronic health record data. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2019, 26, 1083-1090.	2.2	9
69	Blood pressure and body fat percent in women with NMOSD. <i>Brain and Behavior</i> , 2019, 9, e01350.	1.0	9
70	Rapid network meta-analysis using data from Food and Drug Administration approval packages is feasible but with limitations. <i>Journal of Clinical Epidemiology</i> , 2019, 114, 84-94.	2.4	3
71	Two-sample test for correlated data under outcome-dependent sampling with an application to self-reported weight loss data. <i>Statistics in Medicine</i> , 2019, 38, 4999-5009.	0.8	2
72	Plasma Homocysteine Level Is Associated with the Expanded Disability Status Scale in Neuromyelitis Optica Spectrum Disorder. <i>NeuroImmunoModulation</i> , 2019, 26, 258-264.	0.9	6

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73	Association of Race/Ethnicity With Hospital Discharge Disposition After Elective Total Knee Arthroplasty. <i>JAMA Network Open</i> , 2019, 2, e1914259.	2.8	37
74	Investigating safety profiles of human papillomavirus vaccine across group differences using VAERS data and MedDRA. <i>PeerJ</i> , 2019, 7, e7490.	0.9	3
75	Elevated Plasma Homocysteine Levels in Anti-N-methyl-D-aspartate Receptor Encephalitis. <i>Frontiers in Neurology</i> , 2019, 10, 464.	1.1	6
76	Leveraging deep learning to understand health beliefs about the Human Papillomavirus Vaccine from social media. <i>Npj Digital Medicine</i> , 2019, 2, 27.	5.7	22
77	Integration of genetic and clinical information to improve imputation of data missing from electronic health records. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2019, 26, 1056-1063.	2.2	17
78	Determining the Association Between End-of-Life Care Resources and Patient Outcomes in Pennsylvania ICUs*. <i>Critical Care Medicine</i> , 2019, 47, 1591-1598.	0.4	7
79	A Meta-Analysis of the Associations Between the Nurse Work Environment in Hospitals and 4 Sets of Outcomes. <i>Medical Care</i> , 2019, 57, 353-361.	1.1	182
80	Regression analysis of longitudinal data with outcome-dependent sampling and informative censoring. <i>Scandinavian Journal of Statistics</i> , 2019, 46, 831-847.	0.9	7
81	Inflation of type I error rates due to differential misclassification in EHR-derived outcomes: Empirical illustration using breast cancer recurrence. <i>Pharmacoepidemiology and Drug Safety</i> , 2019, 28, 264-268.	0.9	17
82	Automated discovery of test statistics using genetic programming. <i>Genetic Programming and Evolvable Machines</i> , 2019, 20, 127-137.	1.5	3
83	A Bayesian latent class approach for EHR-based phenotyping. <i>Statistics in Medicine</i> , 2019, 38, 74-87.	0.8	19
84	Bayesian hierarchical methods for meta-analysis combining randomized-controlled and single-arm studies. <i>Statistical Methods in Medical Research</i> , 2019, 28, 1293-1310.	0.7	18
85	Difference Between Users and Nonusers of a Patient Portal in Health Behaviors and Outcomes: Retrospective Cohort Study. <i>Journal of Medical Internet Research</i> , 2019, 21, e13146.	2.1	16
86	National Survey of Hospitalists'™ Experiences with Incidental Pulmonary Nodules. <i>Journal of Hospital Medicine</i> , 2019, 14, 353-356.	0.7	6
87	Robust-ODAL: Learning from heterogeneous health systems without sharing patient-level data. , 2019, , .		5
88	ODAL: A one-shot distributed algorithm to perform logistic regressions on electronic health records data from multiple clinical sites. <i>Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing</i> , 2019, 24, 30-41.	0.7	7
89	Automated discovery of test statistics using genetic programming. <i>Genetic Programming and Evolvable Machines</i> , 2019, 20, 127-137.	1.5	0
90	Empirical Comparison of Publication Bias Tests in Meta-Analysis. <i>Journal of General Internal Medicine</i> , 2018, 33, 1260-1267.	1.3	184

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91	A Bayesian hierarchical model for network meta-analysis of multiple diagnostic tests. <i>Biostatistics</i> , 2018, 19, 87-102.	0.9	24
92	PIE: A prior knowledge guided integrated likelihood estimation method for bias reduction in association studies using electronic health records data. <i>Journal of the American Medical Informatics Association: JAMIA</i> , 2018, 25, 345-352.	2.2	15
93	The performance of rapid plasma reagin (RPR) titer in HIV-negative general paresis after neurosyphilis therapy. <i>BMC Infectious Diseases</i> , 2018, 18, 144.	1.3	12
94	Cytosine methylation predicts renal function decline in American Indians. <i>Kidney International</i> , 2018, 93, 1417-1431.	2.6	46
95	Bayesian Mixed Treatment Comparisons Meta-Analysis for Correlated Outcomes Subject to Reporting Bias. <i>Journal of the Royal Statistical Society Series C: Applied Statistics</i> , 2018, 67, 127-144.	0.5	12
96	An improved method for bivariate meta-analysis when within-study correlations are unknown. <i>Research Synthesis Methods</i> , 2018, 9, 73-88.	4.2	11
97	Visualized Emotion Ontology: a model for representing visual cues of emotions. <i>BMC Medical Informatics and Decision Making</i> , 2018, 18, 64.	1.5	14
98	A conditional composite likelihood ratio test with boundary constraints. <i>Biometrika</i> , 2018, 105, 225-232.	1.3	4
99	PSB 2019 Workshop on Text Mining and Visualization for Precision Medicine. , 2018, , .		0
100	Inherited Thrombophilia and the Risk of Arterial Ischemic Stroke: A Systematic Review and Meta-Analysis. <i>Blood</i> , 2018, 132, 2518-2518.	0.6	0
101	ODAL: A one-shot distributed algorithm to perform logistic regressions on electronic health records data from multiple clinical sites. , 2018, , .		13
102	Identification of Rare Adverse Events with Year-varying Reporting Rates for FLU4 Vaccine in VAERS. <i>AMIA ... Annual Symposium proceedings</i> , 2018, 2018, 1544-1551.	0.2	0
103	Neuronal Activity-Induced Sterol Regulatory Element Binding Protein-1 (SREBP1) is Disrupted in Dysbindin-Null Mice—Potential Link to Cognitive Impairment in Schizophrenia. <i>Molecular Neurobiology</i> , 2017, 54, 1699-1709.	1.9	17
104	A composite likelihood method for bivariate meta-analysis in diagnostic systematic reviews. <i>Statistical Methods in Medical Research</i> , 2017, 26, 914-930.	0.7	23
105	Implementing optimal allocation in clinical trials with multiple endpoints. <i>Journal of Statistical Planning and Inference</i> , 2017, 182, 88-99.	0.4	4
106	PLEMT: A Novel Pseudolikelihood-Based EM Test for Homogeneity in Generalized Exponential Tilt Mixture Models. <i>Journal of the American Statistical Association</i> , 2017, 112, 1393-1404.	1.8	7
107	pETM: a penalized Exponential Tilt Model for analysis of correlated high-dimensional DNA methylation data. <i>Bioinformatics</i> , 2017, 33, 1765-1772.	1.8	12
108	A frailty model for recurrent events during alternating restraint and non-restraint time periods. <i>Statistics in Medicine</i> , 2017, 36, 643-654.	0.8	1

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109	Analysis of Individual Differences in Vaccine Pharmacovigilance Using VAERS Data and MedDRA System Organ Classes: A Use Case Study With Trivalent Influenza Vaccine. <i>Biomedical Informatics Insights</i> , 2017, 9, 117822261770062.	4.6	8
110	On meta- and mega-analyses for gene-environment interactions. <i>Genetic Epidemiology</i> , 2017, 41, 876-886.	0.6	2
111	A simple and robust method for multivariate meta-analysis of diagnostic test accuracy. <i>Statistics in Medicine</i> , 2017, 36, 105-121.	0.8	7
112	A Semiparametric Model for VQTL Mapping. <i>Biometrics</i> , 2017, 73, 571-581.	0.8	12
113	A signal detection method for temporal variation of adverse effect with vaccine adverse event reporting system data. <i>BMC Medical Informatics and Decision Making</i> , 2017, 17, 76.	1.5	12
114	Maximum likelihood estimation and EM algorithm of Copas-like selection model for publication bias correction. <i>Biostatistics</i> , 2017, 18, 495-504.	0.9	11
115	Comparing the Human Papillomavirus Vaccination Opinions Trends from Different Twitter User Groups with a Machine Learning Based System and Semiparametric Nonlinear Regression. <i>Studies in Health Technology and Informatics</i> , 2017, 245, 1218.	0.2	2
116	Inference for correlated effect sizes using multiple univariate meta-analyses. <i>Statistics in Medicine</i> , 2016, 35, 1405-1422.	0.8	7
117	Lysosomal iron modulates NMDA receptor-mediated excitation via small GTPase, Dexas1. <i>Molecular Brain</i> , 2016, 9, 38.	1.3	47
118	A meta-analytic framework for detection of genetic interactions. <i>Genetic Epidemiology</i> , 2016, 40, 534-543.	0.6	2
119	Phospholipase C Beta 1: a Candidate Signature Gene for Proneural Subtype High-Grade Glioma. <i>Molecular Neurobiology</i> , 2016, 53, 6511-6525.	1.9	23
120	Meta-analysis of studies with bivariate binary outcomes: a marginal beta-binomial model approach. <i>Statistics in Medicine</i> , 2016, 35, 21-40.	0.8	16
121	A hybrid Bayesian hierarchical model combining cohort and case-control studies for meta-analysis of diagnostic tests: Accounting for partial verification bias. <i>Statistical Methods in Medical Research</i> , 2016, 25, 3015-3037.	0.7	12
122	An Empirical Study for Impacts of Measurement Errors on EHR based Association Studies. <i>AMIA ... Annual Symposium proceedings</i> , 2016, 2016, 1764-1773.	0.2	19
123	An alternative pseudolikelihood method for multivariate random-effects meta-analysis. <i>Statistics in Medicine</i> , 2015, 34, 361-380.	0.8	28
124	A Unification of Models for Meta-Analysis of Diagnostic Accuracy Studies without a Gold Standard. <i>Biometrics</i> , 2015, 71, 538-547.	0.8	27
125	A hybrid model for combining case-control and cohort studies in systematic reviews of diagnostic tests. <i>Journal of the Royal Statistical Society Series C: Applied Statistics</i> , 2015, 64, 469-489.	0.5	14
126	Regression analysis of longitudinal data with irregular and informative observation times. <i>Biostatistics</i> , 2015, 16, 727-739.	0.9	17

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127	Enhancement of Stress Resilience Through Histone Deacetylase 6â€‘Mediated Regulation of Glucocorticoid Receptor Chaperone Dynamics. <i>Biological Psychiatry</i> , 2015, 77, 345-355.	0.7	44
128	Bayesian analysis on meta-analysis of case-control studies accounting for within-study correlation. <i>Statistical Methods in Medical Research</i> , 2015, 24, 836-855.	0.7	23
129	A Class of Pseudolikelihood Ratio Tests for Homogeneity in Exponential Tilt Mixture Models. <i>Scandinavian Journal of Statistics</i> , 2015, 42, 504-517.	0.9	6
130	Semiparametric Tests for Identifying Differentially Methylated Loci With Caseâ€‘Control Designs Using Illumina Arrays. <i>Genetic Epidemiology</i> , 2014, 38, 42-50.	0.6	15
131	Comparing the power of family-based association tests for sequence data with applications in the GAW18 simulated data. <i>BMC Proceedings</i> , 2014, 8, S27.	1.8	3
132	Dexas1, a Small GTPase, Is Required for Glutamate-NMDA Neurotoxicity. <i>Journal of Neuroscience</i> , 2013, 33, 3582-3587.	1.7	60
133	Bivariate random effects models for meta-analysis of comparative studies with binary outcomes: Methods for the absolute risk difference and relative risk. <i>Statistical Methods in Medical Research</i> , 2012, 21, 621-633.	0.7	58
134	On the asymptotic behaviour of the pseudolikelihood ratio test statistic with boundary problems. <i>Biometrika</i> , 2010, 97, 603-620.	1.3	33