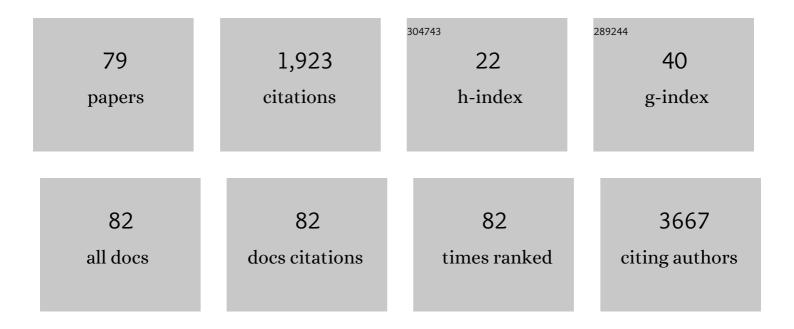
Jinbo Chen

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A Robust Approach for Electronic Health Record–Based Case-Control Studies with Contaminated Case Pools. Biometrics, 2023, 79, 2023-2035.	1.4	0
2	Impact of Behavioral Nudges on the Quality of Serious Illness Conversations Among Patients With Cancer: Secondary Analysis of a Randomized Controlled Trial. JCO Oncology Practice, 2022, 18, e495-e503.	2.9	3
3	A systematic review of the natural history and biomarkers of primary lecithin:cholesterol acyltransferase deficiency. Journal of Lipid Research, 2022, 63, 100169.	4.2	8
4	Validation of Breast Cancer Risk Models by Race/Ethnicity, Family History and Molecular Subtypes. Cancers, 2022, 14, 45.	3.7	11
5	Case contamination in electronic health records based caseâ€control studies. Biometrics, 2021, 77, 67-77.	1.4	5
6	Exome-wide evaluation of rare coding variants using electronic health records identifies new gene–phenotype associations. Nature Medicine, 2021, 27, 66-72.	30.7	44
7	Testing calibration of phenotyping models using positive-only electronic health record data. Biostatistics, 2021, , .	1.5	1
8	Respiratory Motion Mitigation and Repeatability of Two Diffusion-Weighted MRI Methods Applied to a Murine Model of Spontaneous Pancreatic Cancer. Tomography, 2021, 7, 66-79.	1.8	3
9	Risk factors for an advanced breast cancer diagnosis within 2 years of a negative mammogram. Cancer, 2021, 127, 3334-3342.	4.1	9
10	Trajectories of mortality risk among patients with cancer and associated end-of-life utilization. Npj Digital Medicine, 2021, 4, 104.	10.9	1
11	Relationship of established risk factors with breast cancer subtypes. Cancer Medicine, 2021, 10, 6456-6467.	2.8	45
12	Covariate adjusted inference of parentâ€ofâ€origin effects using case–control mother–child paired multilocus genotype data. Genetic Epidemiology, 2021, 45, 830-847.	1.3	2
13	Two-Stage Approaches to Accounting for Patient Heterogeneity in Machine Learning Risk Prediction Models in Oncology. JCO Clinical Cancer Informatics, 2021, 5, 1015-1023.	2.1	1
14	Maternal Proinflammatory Adipokines Throughout Pregnancy and Neonatal Size and Body Composition: A Prospective Study. Current Developments in Nutrition, 2021, 5, nzab113.	0.3	5
15	A Multi-Marker Test for Analyzing Paired Genetic Data in Transplantation. Frontiers in Genetics, 2021, 12, 745773.	2.3	2
16	OUP accepted manuscript. Biostatistics, 2021, , .	1.5	0
17	Extreme-value sampling design is cost-beneficial only with a valid statistical approach for exposure–secondary outcome association analyses. Statistical Methods in Medical Research, 2020, 29, 466-480.	1.5	2
18	A Population-Level Analysis of Pituitary Carcinoma from the National Cancer Database. Journal of Neurological Surgery, Part B: Skull Base, 2020, 81, 180-186.	0.8	6

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19	Novel twoâ€phase sampling designs for studying binary outcomes. Biometrics, 2020, 76, 210-223.	1.4	4
20	A maximum likelihood approach to electronic health record phenotyping using positive and unlabeled patients. Journal of the American Medical Informatics Association: JAMIA, 2020, 27, 119-126.	4.4	13
21	Precision prophylaxis: Identifying the optimal timing for risk-reducing salpingo-oophorectomy based on type of BRCA1 and BRCA2 cluster region mutations. Gynecologic Oncology, 2020, 156, 363-376.	1.4	3
22	Adipokines in early and mid-pregnancy and subsequent risk of gestational diabetes: a longitudinal study in a multiracial cohort. BMJ Open Diabetes Research and Care, 2020, 8, e001333.	2.8	26
23	Joint testing of donor and recipient genetic matching scores and recipient genotype has robust power for finding genes associated with transplant outcomes. Genetic Epidemiology, 2020, 44, 893-907.	1.3	7
24	Validation of a Machine Learning Algorithm to Predict 180-Day Mortality for Outpatients With Cancer. JAMA Oncology, 2020, 6, 1723.	7.1	71
25	Phenotyping issues for exploring electronic health records to design clinical trials. Clinical Trials, 2020, 17, 402-404.	1.6	0
26	Phenome-wide association analysis suggests the APOL1 linked disease spectrum primarily drives kidney-specific pathways. Kidney International, 2020, 97, 1032-1041.	5.2	20
27	Penetrance of Breast and Ovarian Cancer in Women Who Carry a BRCA1/2 Mutation and Do Not Use Risk-Reducing Salpingo-Oophorectomy: An Updated Meta-Analysis. JNCI Cancer Spectrum, 2020, 4, pkaa029.	2.9	41
28	An efficient and computationally robust statistical method for analyzing case-control mother–offspring pair genetic association studies. Annals of Applied Statistics, 2020, 14, .	1.1	4
29	Genome-wide association study of peripheral artery disease in the Million Veteran Program. Nature Medicine, 2019, 25, 1274-1279.	30.7	177
30	Impact of payer status on survival in parotid malignancy. American Journal of Otolaryngology - Head and Neck Medicine and Surgery, 2019, 40, 555-559.	1.3	7
31	High risk of metabolic syndrome after delivery in pregnancies complicated by gestational diabetes. Diabetes Research and Clinical Practice, 2019, 150, 219-226.	2.8	31
32	Hearing and Quality of Life Over Time in Vestibular Schwannoma Patients: Observation Compared to Stereotactic Radiosurgery. Otology and Neurotology, 2019, 40, 1094-1100.	1.3	12
33	Maternal adipokines longitudinally measured across pregnancy and their associations with neonatal size, length, and adiposity. International Journal of Obesity, 2019, 43, 1422-1434.	3.4	31
34	A Population-Level Analysis of Pituitary Carcinoma from the National Cancer Database. Journal of Neurological Surgery, Part B: Skull Base, 2019, 80, .	0.8	0
35	Ageing perceptions and non-adherence to aromatase inhibitors among breast cancer survivors. European Journal of Cancer, 2018, 91, 145-152.	2.8	20
36	Mucoepidermoid carcinoma of the parotid gland: A National Cancer Database study. American Journal of Otolaryngology - Head and Neck Medicine and Surgery, 2018, 39, 321-326.	1.3	40

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37	Safety of outpatient thyroidectomy: Review of the American College of Surgeons National Surgical Quality Improvement Program. Laryngoscope, 2018, 128, 1249-1254.	2.0	32
38	Clinical Factors Associated With Reoperation and Prolonged Length of Stay in Free Tissue Transfer to Oncologic Head and Neck Defects. JAMA Facial Plastic Surgery, 2018, 20, 154-159.	2.1	32
39	Clinicopathologic Factors Predictive of Occult Lymph Node Involvement in Cutaneous Head and Neck Melanoma. Otolaryngology - Head and Neck Surgery, 2018, 158, 489-496.	1.9	8
40	AzBio Speech Understanding Performance in Quiet and Noise in High Performing Cochlear Implant Users. Otology and Neurotology, 2018, 39, 571-575.	1.3	16
41	Breast and ovarian cancer penetrance of <i>BRCA1/2</i> mutations among Hong Kong women. Oncotarget, 2018, 9, 25025-25033.	1.8	8
42	Genomic Risk Stratification Predicts All-Cause Mortality After Cardiac Catheterization. Circulation Genomic and Precision Medicine, 2018, 11, e002352.	3.6	16
43	Adjustment of nonconfounding covariates in case-control genetic association studies. Annals of Applied Statistics, 2018, 12, .	1.1	5
44	Complications Associated with Mortality after Head and Neck Surgery: An Analysis of the NSQIP Database. Otolaryngology - Head and Neck Surgery, 2017, 156, 504-510.	1.9	11
45	Association Between Patient-Reported Medication Adherence and Anticoagulation Control. American Journal of Medicine, 2017, 130, 1092-1098.e2.	1.5	15
46	Phenotype validation in electronic health records based genetic association studies. Genetic Epidemiology, 2017, 41, 790-800.	1.3	8
47	Patient, disease, and treatment factors associated with overall survival in esthesioneuroblastoma. International Forum of Allergy and Rhinology, 2017, 7, 1186-1194.	2.8	33
48	Lipoprotein(a) and Risk of Myocardial Infarction and Death in Chronic Kidney Disease. Arteriosclerosis, Thrombosis, and Vascular Biology, 2017, 37, 1971-1978.	2.4	44
49	Influence of common and rare genetic variation on warfarin dose among African–Americans and European–Americans using the exome array. Pharmacogenomics, 2017, 18, 1059-1073.	1.3	12
50	Perceived barriers to treatment predict adherence to aromatase inhibitors among breast cancer survivors. Cancer, 2017, 123, 169-176.	4.1	39
51	Association between Breast Parenchymal Complexity and False-Positive Recall From Digital Mammography Versus Breast Tomosynthesis. Academic Radiology, 2016, 23, 977-986.	2.5	4
52	Efficient unified rare variant association test by modeling the population genetic distribution in case ontrol studies. Genetic Epidemiology, 2016, 40, 579-590.	1.3	2
53	Using family members to augment genetic case–control studies of a lifeâ€ŧhreatening disease. Statistics in Medicine, 2016, 35, 2815-2830.	1.6	4
54	Self-renewal of CD133hi cells by IL6/Notch3 signalling regulates endocrine resistance in metastatic breast cancer. Nature Communications, 2016, 7, 10442.	12.8	144

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55	Quantitative assessment of background parenchymal enhancement in breast MRI predicts response to risk-reducing salpingo-oophorectomy: preliminary evaluation in a cohort of BRCA1/2 mutation carriers. Breast Cancer Research, 2015, 17, 67.	5.0	49
56	Arthralgia among women taking aromatase inhibitors: is there a shared inflammatory mechanism with co-morbid fatigue and insomnia?. Breast Cancer Research, 2015, 17, 89.	5.0	35
57	Retrospective Likelihood-Based Methods for Analyzing Case-Cohort Genetic Association Studies. Biometrics, 2015, 71, 960-968.	1.4	3
58	Parent-of-Origin Effects of the APOB Gene on Adiposity in Young Adults. PLoS Genetics, 2015, 11, e1005573.	3.5	16
59	Parenchymal texture analysis in digital mammography: robust texture feature identification and equivalence across devices. Journal of Medical Imaging, 2015, 2, 024501.	1.5	19
60	A robust association test for detecting genetic variants with heterogeneous effects. Biostatistics, 2015, 16, 5-16.	1.5	1
61	Associations between breast density and a panel of single nucleotide polymorphisms linked to breast cancer risk: a cohort study with digital mammography. BMC Cancer, 2015, 15, 143.	2.6	15
62	The use of the Gail model, body mass index and SNPs to predict breast cancer among women with abnormal (BI-RADS 4) mammograms. Breast Cancer Research, 2015, 17, 1.	5.0	124
63	RE: Breast Cancer Risk After Salpingo-Oophorectomy in Healthy BRCA1/2 Mutation Carriers: Revisiting the Evidence for Risk Reduction. Journal of the National Cancer Institute, 2015, 107, .	6.3	23
64	Preliminary evaluation of the publicly available Laboratory for Breast Radiodensity Assessment (LIBRA) software tool: comparison of fully automated area and volumetric density measures in a case–control study with digital mammography. Breast Cancer Research, 2015, 17, 117.	5.0	68
65	Breast density and parenchymal texture measures as potential risk factors for estrogen-receptor positive breast cancer. Proceedings of SPIE, 2014, 9035, 90351D.	0.8	12
66	A robust test for quantitative trait analysis with model uncertainty in genetic association studies. Statistics and Its Interface, 2014, 7, 61-68.	0.3	2
67	A Multiâ€Locus Likelihood Method for Assessing Parentâ€ofâ€Origin Effects Using Caseâ€Control Motherâ€Child Pairs. Genetic Epidemiology, 2013, 37, 152-162.	1.3	9
68	Efficient designs of gene–environment interaction studies: implications of Hardy–Weinberg equilibrium and gene–environment independence. Statistics in Medicine, 2012, 31, 2516-2530.	1.6	12
69	Semiparametric Maximum Likelihood Methods for Analyzing Genetic and Environmental Effects with Caseâ€Control Mother–Child Pair Data. Biometrics, 2012, 68, 869-877.	1.4	10
70	Likelihood ratio tests for maternal and fetal genetic effects on obstetric complications. Genetic Epidemiology, 2009, 33, 526-538.	1.3	14
71	Testing Hardyâ€Weinberg equilibrium using motherâ€child caseâ€control samples. Genetic Epidemiology, 2009, 33, 539-548.	1.3	1
72	Statistical Methods for Analyzing Two-Phase Studies. Frontiers of Statistics, 2009, , 127-155.	0.2	0

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73	Breast Cancer Relative Hazard Estimates From Case–Control and Cohort Designs With Missing Data on Mammographic Density. Journal of the American Statistical Association, 2008, 103, 976-988.	3.1	6
74	Exploiting Hardy-Weinberg Equilibrium for Efficient Screening of Single SNP Associations from Case-Control Studies. Human Heredity, 2007, 63, 196-204.	0.8	43
75	A partially linear tree-based regression model for assessing complex joint gene–gene and gene–environment effects. Genetic Epidemiology, 2007, 31, 238-251.	1.3	22
76	Conditional Likelihood Methods for Haplotype-Based Association Analysis Using Matched Case-Control Data. Biometrics, 2007, 63, 1099-1107.	1.4	2
77	Haplotype-Based Association Analysis in Cohort and Nested Case-Control Studies. Biometrics, 2006, 62, 28-35.	1.4	10
78	Projecting Absolute Invasive Breast Cancer Risk in White Women With a Model That Includes Mammographic Density. Journal of the National Cancer Institute, 2006, 98, 1215-1226.	6.3	317
79	A Haplotype-Based Test of Association Using Data from Cohort and Nested Case-Control Epidemiologic Studies. Human Heredity, 2004, 58, 18-29.	0.8	18