## **Chixiang Chen**

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A Robust Consistent Information Criterion for Model Selection Based on Empirical Likelihood. Statistica Sinica, 2022, , .	0.3	1
2	Improving main analysis by borrowing information fromÂauxiliary data. Statistics in Medicine, 2022, 41, 567-579.	1.6	5
3	A statistical framework for recovering pseudo-dynamic networks from static data. Bioinformatics, 2022, , .	4.1	2
4	Semiparametric marginal methods for clustered data adjusting for informative cluster size with nonignorable zeros. Biometrical Journal, 2022, 64, 898-911.	1.0	3
5	A multiple robust propensity score method for longitudinal analysis with intermittent missing data. Biometrics, 2021, 77, 519-532.	1.4	14
6	Joint modeling of longitudinal data with informative cluster size adjusted for zeroâ€inflation and a dependent terminal event. Statistics in Medicine, 2021, 40, 4582-4596.	1.6	3
7	A Computational Atlas of Tissue-specific Regulatory Networks. Frontiers in Systems Biology, 2021, 1, .	0.7	3
8	Assessing predictive accuracy of survival regressions subject to nonindependent censoring. Statistics in Medicine, 2020, 39, 469-480.	1.6	1
9	Prostate cancer in Pennsylvania: The role of older age at diagnosis, aggressiveness, and environmental risk factors on treatment and mortality using data from the Pennsylvania Cancer Registry. Cancer Medicine, 2020, 9, 3623-3633.	2.8	8
10	Association of SNP–SNP Interactions of Surfactant Protein Genes with Pediatric Acute Respiratory Failure. Journal of Clinical Medicine, 2020, 9, 1183.	2.4	13
11	SNP and Haplotype Interaction Models Reveal Association of Surfactant Protein Gene Polymorphisms With Hypersensitivity Pneumonitis of Mexican Population. Frontiers in Medicine, 2020, 7, 588404.	2.6	14
12	Greater Exercise Pressor Response Is Associated With Impaired Claudication Outcomes in Symptomatic Peripheral Artery Disease. Angiology, 2019, 70, 220-228.	1.8	6
13	An omnidirectional visualization model of personalized gene regulatory networks. Npj Systems Biology and Applications, 2019, 5, 38.	3.0	20
14	Empirical-Likelihood-Based Criteria for Model Selection on Marginal Analysis of Longitudinal Data With Dropout Missingness. Biometrics, 2019, 75, 950-965.	1.4	8
15	Simultaneous confidence bands for growth incidence curves in weighted sup-norm metrics. Statistics, 2019, 53, 1269-1288.	0.6	1
16	A unified DNA sequence and non―DNA sequence mapping model of complex traits. Plant Journal, 2019, 99, 784-795.	5.7	2
17	<i>np</i> <sup>2</sup> <scp>QTL</scp> : networking phenotypic plasticity quantitative trait loci across heterogeneous environments. Plant Journal, 2019, 99, 796-806.	5.7	7
18	Vascular Inflammation, Calf Muscle Oxygen Saturation, and Blood Glucose are Associated With Exercise Pressor Response in Symptomatic Peripheral Artery Disease. Angiology, 2019, 70, 747-755.	1.8	6

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19	Inhibition of UDP-glucose dehydrogenase by 6-thiopurine and its oxidative metabolites: Possible mechanism for its interaction within the bilirubin excretion pathway and 6TP associated liver toxicity. Journal of Pharmaceutical and Biomedical Analysis, 2018, 151, 106-115.	2.8	4
20	Effect of adipose tissue thickness, muscle site, and sex on near-infrared spectroscopy derived total-[hemoglobin + myoglobin]. Journal of Applied Physiology, 2017, 123, 1571-1578.	2.5	48
21	Single Nucleotide Polymorphisms (SNP) and SNP-SNP Interactions of the Surfactant Protein Genes Are Associated With Idiopathic Pulmonary Fibrosis in a Mexican Study Group; Comparison With Hypersensitivity Pneumonitis. Frontiers in Immunology, 0, 13, .	4.8	2