

# Tao Lu

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

29  
papers

233  
citations

6  
h-index

15  
g-index

29  
ext. papers

266  
ext. citations

1.7  
avg, IF

3.36  
L-index

#	Paper	IF	Citations
29	Estimation of the error distribution function for partial linear single-index models. <i>Communications in Statistics Part B: Simulation and Computation</i> , <b>2020</b> , 49, 29-44	0.6	3
28	Multiplicative regression models with distortion measurement errors. <i>Statistical Papers</i> , <b>2020</b> , 61, 2031-2057		14
27	Bayesian panel smooth transition model with spatial correlation. <i>PLoS ONE</i> , <b>2019</b> , 14, e0211467	3.7	1
26	Association between depressive symptoms and poor sleep quality among Han and Manchu ethnicities in a large, rural, Chinese population. <i>PLoS ONE</i> , <b>2019</b> , 14, e0226562	3.7	5
25	Novel polymeric biomaterial poly(butyl-2-cyanoacrylate) nanowires: synthesis, characterization and formation mechanism. <i>Colloids and Surfaces B: Biointerfaces</i> , <b>2019</b> , 175, 454-462	6	4
24	Partially linear mixed-effects joint models for skewed and missing longitudinal competing risks outcomes. <i>Journal of Biopharmaceutical Statistics</i> , <b>2019</b> , 29, 971-989	1.3	2
23	Testing symmetry based on empirical likelihood. <i>Journal of Applied Statistics</i> , <b>2018</b> , 45, 2429-2454	1	9
22	Mixed-effects location and scale Tobit joint models for heterogeneous longitudinal data with skewness, detection limits, and measurement errors. <i>Statistical Methods in Medical Research</i> , <b>2018</b> , 27, 3525-3543	2.3	2
21	Design of experiment for nonlinear dynamic gene regulatory network identification. <i>Journal of Biopharmaceutical Statistics</i> , <b>2018</b> , 28, 402-412	1.3	
20	Objective Bayesian hypothesis testing and estimation for the intraclass model. <i>Statistical Theory and Related Fields</i> , <b>2018</b> , 2, 37-47	0.3	
19	Bayesian inference on mixed-effects varying-coefficient joint models with skew- t distribution for longitudinal data with multiple features. <i>Statistical Methods in Medical Research</i> , <b>2017</b> , 26, 1146-1164	2.3	3
18	Bayesian nonparametric mixed-effects joint model for longitudinal-competing risks data analysis in presence of multiple data features. <i>Statistical Methods in Medical Research</i> , <b>2017</b> , 26, 2407-2423	2.3	3
17	Simultaneous inference for semiparametric mixed-effects joint models with skew distribution and covariate measurement error for longitudinal competing risks data analysis. <i>Journal of Biopharmaceutical Statistics</i> , <b>2017</b> , 27, 1009-1027	1.3	2
16	Modeling Longitudinal-Competing Risks Data With Skew Distribution and Mismeasured Covariate. <i>Statistics in Biopharmaceutical Research</i> , <b>2017</b> , 9, 73-84	1.2	
15	Bayesian inference on mixed-effects location scale models with skew-t distribution and mismeasured covariates for longitudinal data. <i>Statistics in Medicine</i> , <b>2017</b> , 36, 2614-2629	2.3	1
14	Bayesian varying coefficient mixed-effects joint models with asymmetry and missingness. <i>Statistical Modelling</i> , <b>2017</b> , 17, 117-141	0.7	1
13	Jointly modeling skew longitudinal survival data with missingness and mismeasured covariates. <i>Journal of Applied Statistics</i> , <b>2017</b> , 44, 2354-2367	1	3

12	Bayesian inference on longitudinal-survival data with multiple features. <i>Computational Statistics</i> , <b>2017</b> , 32, 845-866	1	2
11	Bayesian inference on partially linear mixed-effects joint models for longitudinal data with multiple features. <i>Computational Statistics</i> , <b>2017</b> , 32, 179-196	1	3
10	Bayesian semiparametric mixed-effects joint models for analysis of longitudinal-competing risks data with skew distribution. <i>Statistics and Its Interface</i> , <b>2017</b> , 10, 441-450	0.4	1
9	Attitudes towards suicide in urban and rural China: a population based, cross-sectional study. <i>BMC Psychiatry</i> , <b>2016</b> , 16, 162	4.2	58
8	Investigate Data Dependency for Dynamic Gene Regulatory Network Identification through High-dimensional Differential Equation Approach. <i>Communications in Statistics Part B: Simulation and Computation</i> , <b>2016</b> , 45, 2377-2391	0.6	1
7	Skew-t partially linear mixed-effects models for AIDS clinical studies. <i>Journal of Biopharmaceutical Statistics</i> , <b>2016</b> , 26, 899-911	1.3	
6	Mixed-effects varying-coefficient model with skewed distribution coupled with cause-specific varying-coefficient hazard model with random-effects for longitudinal-competing risks data analysis. <i>Journal of Biopharmaceutical Statistics</i> , <b>2016</b> , 26, 519-33	1.3	2
5	Robust variable selection method for nonparametric differential equation models with application to nonlinear dynamic gene regulatory network analysis. <i>Journal of Biopharmaceutical Statistics</i> , <b>2016</b> , 26, 712-24	1.3	
4	Bayesian structured variable selection in linear regression models. <i>Computational Statistics</i> , <b>2015</b> , 30, 205-229	1	6
3	Sparse Additive Ordinary Differential Equations for Dynamic Gene Regulatory Network Modeling. <i>Journal of the American Statistical Association</i> , <b>2014</b> , 109, 700-716	2.8	53
2	A refined parameter estimating approach for HIV dynamic model. <i>Journal of Applied Statistics</i> , <b>2014</b> , 41, 1645-1657	1	1
1	High Dimensional ODEs Coupled with Mixed-Effects Modeling Techniques for Dynamic Gene Regulatory Network Identification. <i>Journal of the American Statistical Association</i> , <b>2011</b> , 106, 1242-1258	2.8	53