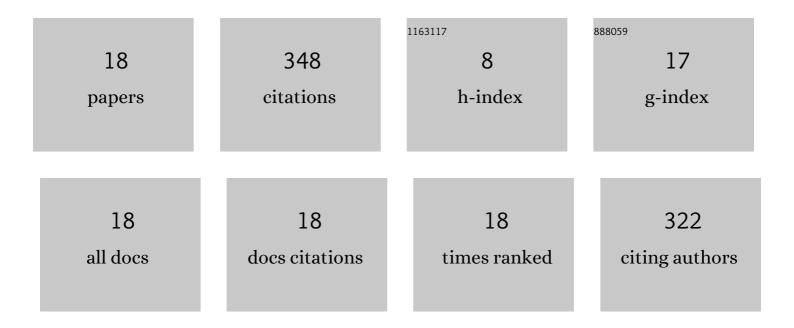
## Longhai Li

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3867904/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	A New Regression Model: Modal Linear Regression. Scandinavian Journal of Statistics, 2014, 41, 656-671.	1.4	120
2	Deep learning for brain disorder diagnosis based on fMRI images. Neurocomputing, 2022, 469, 332-345.	5.9	51
3	A comparison of residual diagnosis tools for diagnosing regression models for count data. BMC Medical Research Methodology, 2020, 20, 175.	3.1	38
4	Sleep duration and sleep disturbances in association with falls among the middle-aged and older adults in China: a population-based nationwide study. BMC Geriatrics, 2018, 18, 196.	2.7	36
5	Antimicrobial resistance genetic factor identification from whole-genome sequence data using deep feature selection. BMC Bioinformatics, 2019, 20, 535.	2.6	22
6	Predictive analysis methods for human microbiome data with application to Parkinson's disease. PLoS ONE, 2020, 15, e0237779.	2.5	21
7	Approximating cross-validatory predictive evaluation in Bayesian latent variable models with integrated IS and WAIC. Statistics and Computing, 2016, 26, 881-897.	1.5	19
8	Bayesian Hyper-LASSO Classification for Feature Selection with Application to Endometrial Cancer RNA-seq Data. Scientific Reports, 2020, 10, 9747.	3.3	10
9	Fully Bayesian logistic regression with hyper-LASSO priors for high-dimensional feature selection. Journal of Statistical Computation and Simulation, 2018, 88, 2827-2851.	1.2	9
10	Model diagnostics for censored regression via randomized survival probabilities. Statistics in Medicine, 2021, 40, 1482-1497.	1.6	6
11	Compressing parameters in Bayesian high-order models with application to logistic sequence models. Bayesian Analysis, 2008, 3, .	3.0	3
12	Bias-Corrected Hierarchical Bayesian Classification With a Selected Subset of High-Dimensional Features. Journal of the American Statistical Association, 2012, 107, 120-134.	3.1	3
13	An online Bayesian mixture labelling method by minimizing deviance of classification probabilities to reference labels. Journal of Statistical Computation and Simulation, 2014, 84, 310-323.	1.2	3
14	Are Bayesian Inferences Weak for Wasserman's Example?. Communications in Statistics Part B: Simulation and Computation, 2010, 39, 655-667.	1.2	2
15	Computational identification of harmful mutation regions to the activity of transposable elements. BMC Genomics, 2017, 18, 862.	2.8	2
16	Randomized quantile residuals for diagnosing zero-inflated generalized linear mixed models with applications to microbiome count data. BMC Bioinformatics, 2021, 22, 564.	2.6	2
17	Estimating crossâ€validatory predictive <i>p</i> â€values with integrated importance sampling for disease mapping models. Statistics in Medicine, 2017, 36, 2220-2236.	1.6	1
18	Impact of misspecified residual correlation structure on the parameter estimates in a shared spatial frailty model. Journal of Statistical Computation and Simulation, 2017, 87, 2384-2410.	1.2	0