Jing Lei

List of Publications by Year in descending order

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516710 610901 1,840 27 16 24 citations h-index g-index papers 28 28 28 2064 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Bias-Adjusted Spectral Clustering in Multi-Layer Stochastic Block Models. Journal of the American Statistical Association, 2023, 118, 2433-2445.	3.1	9
2	Model selection properties of forward selection and sequential crossâ€validation for highâ€dimensional regression. Canadian Journal of Statistics, 2022, 50, 454-470.	0.9	4
3	Consistent estimation of the number of communities in stochastic block models using crossâ€validation. Stat, 2022, 11, e426.	0.4	O
4	Exponential-Family Embedding With Application to Cell Developmental Trajectories for Single-Cell RNA-Seq Data. Journal of the American Statistical Association, 2021, 116, 457-470.	3.1	7
5	Identification of cell-type-specific marker genes from co-expression patterns in tissue samples. Bioinformatics, 2021, 37, 3228-3234.	4.1	9
6	Network representation using graph root distributions. Annals of Statistics, 2021, 49, .	2.6	3
7	Rejoinder for "Exponential-Family Embedding With Application to Cell Developmental Trajectories for Single-Cell RNA-Seq Data― Journal of the American Statistical Association, 2021, 116, 478-480.	3.1	0
8	Cross-Validation With Confidence. Journal of the American Statistical Association, 2020, 115, 1978-1997.	3.1	37
9	Consistent community detection in multi-layer network data. Biometrika, 2020, 107, 61-73.	2.4	26
10	Least Ambiguous Set-Valued Classifiers With Bounded Error Levels. Journal of the American Statistical Association, 2019, 114, 223-234.	3.1	36
11	Squared-norm empirical processes. Statistics and Probability Letters, 2019, 150, 108-113.	0.7	1
12	Semisoft clustering of single-cell data. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 466-471.	7.1	71
13	Network Cross-Validation for Determining the Number of Communities in Network Data. Journal of the American Statistical Association, 2018, 113, 241-251.	3.1	82
14	Distribution-Free Predictive Inference for Regression. Journal of the American Statistical Association, 2018, 113, 1094-1111.	3.1	246
15	Differentially Private Model Selection with Penalized and Constrained Likelihood. Journal of the Royal Statistical Society Series A: Statistics in Society, 2018, 181, 609-633.	1.1	6
16	Sparse subspace linear discriminant analysis. Statistics, 2018, 52, 782-800.	0.6	1
17	Testing high-dimensional covariance matrices, with application to detecting schizophrenia risk genes. Annals of Applied Statistics, 2017, 11, 1810-1831.	1.1	20
18	A goodness-of-fit test for stochastic block models. Annals of Statistics, 2016, 44, .	2.6	82

#	Article	IF	CITATIONS
19	Network assisted analysis to reveal the genetic basis of autism. Annals of Applied Statistics, 2015, 9, 1571-1600.	1.1	43
20	Consistency of spectral clustering in stochastic block models. Annals of Statistics, 2015, 43, .	2.6	249
21	Sparsistency and agnostic inference in sparse PCA. Annals of Statistics, 2015, 43, .	2.6	24
22	The autism-associated chromatin modifier CHD8 regulates other autism risk genes during human neurodevelopment. Nature Communications, 2015, 6, 6404.	12.8	316
23	A conformal prediction approach to explore functional data. Annals of Mathematics and Artificial Intelligence, 2015, 74, 29-43.	1.3	29
24	Distribution-free Prediction Bands for Non-parametric Regression. Journal of the Royal Statistical Society Series B: Statistical Methodology, 2014, 76, 71-96.	2.2	80
25	Distribution-Free Prediction Sets. Journal of the American Statistical Association, 2013, 108, 278-287.	3.1	76
26	Minimax sparse principal subspace estimation in high dimensions. Annals of Statistics, 2013, 41, .	2.6	86
27	Differential privacy and robust statistics. , 2009, , .		297