

Ultrahigh dimensional feature selection: beyond the lin

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Stability Selection. Journal of the Royal Statistical Society Series B: Statistical Methodology, 2010, 72, 417-473.	1.1	1,578
2	Model-Free Feature Screening for Ultrahigh-Dimensional Data. Journal of the American Statistical Association, 2011, 106, 1464-1475.	1.8	357
3	Detection of rare functional variants using group ISIS. BMC Proceedings, 2011, 5, S108.	1.8	5
4	A variable selection method for genome-wide association studies. Bioinformatics, 2011, 27, 1-8.	1.8	132
5	Feature Screening via Distance Correlation Learning. Journal of the American Statistical Association, 2012, 107, 1129-1139.	1.8	510
6	A comparison between subset selection and L1 regularisation with an application in spectroscopy. Chemometrics and Intelligent Laboratory Systems, 2012, 118, 127-138.	1.8	20
8	An Enduring Interest in Classification: Supervised and Unsupervised. , 2012, , 147-171.		0
9	Variance Estimation Using Refitted Cross-Validation in Ultrahigh Dimensional Regression. Journal of the Royal Statistical Society Series B: Statistical Methodology, 2012, 74, 37-65.	1.1	166
10	Focused estimation and model averaging with penalization methods: an overview. Statistica Neerlandica, 2012, 66, 272-287.	0.9	14
11	Feature selection in corporate credit rating prediction. Knowledge-Based Systems, 2013, 51, 72-84.	4.0	87
12	Independent Screening for Single-Index Hazard rate Models with Ultrahigh Dimensional Features. Journal of the Royal Statistical Society Series B: Statistical Methodology, 2013, 75, 217-245.	1.1	66
13	Statistical analysis of big data on pharmacogenomics. Advanced Drug Delivery Reviews, 2013, 65, 987-1000.	6.6	39
14	Correlated Component Regression: Re-thinking Regression in the Presence of Near Collinearity. Springer Proceedings in Mathematics and Statistics, 2013, , 65-78.	0.1	17
15	Feature Screening for Ultrahigh Dimensional Categorical Data With Applications. Journal of Business and Economic Statistics, 2014, 32, 237-244.	1.8	51
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20	Challenges of Big Data analysis. National Science Review, 2014, 1, 293-314.	4.6	954
21	A feature selection method for classification within functional genomics experiments based on the proportional overlapping score. BMC Bioinformatics, 2014, 15, 274.	1.2	40
22	Combining techniques for screening and evaluating interaction terms on high-dimensional time-to-event data. BMC Bioinformatics, 2014, 15, 58.	1.2	6
23	On selecting interacting features from high-dimensional data. Computational Statistics and Data Analysis, 2014, 71, 694-708.	0.7	23
24	A forest-based feature screening approach for large-scale genome data with complex structures. BMC Genetics, 2015, 16, 148.	2.7	4
25	Comment. Journal of the American Statistical Association, 2015, 110, 1439-1442.	1.8	1
26	Big Data in Survey Research. Public Opinion Quarterly, 2015, 79, 839-880.	0.9	115
27	Model-Free Feature Screening for Ultrahigh Dimensional Discriminant Analysis. Journal of the American Statistical Association, 2015, 110, 630-641.	1.8	161
28	Towards scalable fuzzy-rough feature selection. Information Sciences, 2015, 323, 1-15.	4.0	49
29	A Bayesian Approach to Sparse Cox Regression in High-Dimensional Survival Analysis. Lecture Notes in Computer Science, 2015, , 425-437.	1.0	4
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33	Feature screening in ultrahigh dimensional cox's model. Statistica Sinica, 2016, 26, 881-901.	0.2	21
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37	Model-free sure screening via maximum correlation. Journal of Multivariate Analysis, 2016, 148, 89-106.	0.5	8

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39	A data-driven approach to conditional screening of high-dimensional variables. Stat, 2016, 5, 200-212.	0.3	11
40	Local independence feature screening for nonparametric and semiparametric models by marginal empirical likelihood. Annals of Statistics, 2016, 44, 515-539.	1.4	28
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49	PLEMT: A Novel Pseudolikelihood-Based EM Test for Homogeneity in Generalized Exponential Tilt Mixture Models. Journal of the American Statistical Association, 2017, 112, 1393-1404.	1.8	7
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51	TSGSIS: a high-dimensional grouped variable selection approach for detection of whole-genome SNP-SNP interactions. Bioinformatics, 2017, 33, 3595-3602.	1.8	5
52	A Robust Model-Free Feature Screening Method for Ultrahigh-Dimensional Data. Journal of Computational and Graphical Statistics, 2017, 26, 803-813.	0.9	15
53	Improvement Screening for Ultra-High Dimensional Data with Censored Survival Outcomes and Varying Coefficients. International Journal of Biostatistics, 2017, 13, .	0.4	8
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57	Nearly assumptionless screening for the mutually-exciting multivariate Hawkes process. <i>Electronic Journal of Statistics</i> , 2017, 11, 1207-1234.	0.4	7
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60	An adaptive threshold determination method of feature screening for genomic selection. <i>BMC Bioinformatics</i> , 2017, 18, 212.	1.2	2
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62	Model free feature screening for ultrahigh dimensional data with responses missing at random. <i>Computational Statistics and Data Analysis</i> , 2017, 105, 201-216.	0.7	30
63	High-dimensional variable selection in regression and classification with missing data. <i>Signal Processing</i> , 2017, 131, 1-7.	2.1	7
64	A robust variable screening method for high-dimensional data. <i>Journal of Applied Statistics</i> , 2017, 44, 1839-1855.	0.6	6
65	Feature selection of ultrahigh-dimensional covariates with survival outcomes: a selective review. <i>Applied Mathematics</i> , 2017, 32, 379-396.	0.6	11
66	Partition-based ultrahigh-dimensional variable screening. <i>Biometrika</i> , 2017, 104, 785-800.	1.3	15
67	Model-free conditional feature screening for ultra-high dimensional right censored data. <i>Journal of Statistical Computation and Simulation</i> , 2018, 88, 2425-2446.	0.7	4
68	Statistical testing and power analysis for brain-wide association study. <i>Medical Image Analysis</i> , 2018, 47, 15-30.	7.0	18
69	Model-free feature screening for high-dimensional survival data. <i>Science China Mathematics</i> , 2018, 61, 1617-1636.	0.8	8
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72	Nonparametric independence feature screening for ultrahigh-dimensional survival data. <i>Metrika</i> , 2018, 81, 821-847.	0.5	1
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76	Robust feature screening for ultra-high dimensional right censored data via distance correlation. <i>Computational Statistics and Data Analysis</i> , 2018, 119, 118-138.	0.7	27
77	Variable screening for ultrahigh dimensional heterogeneous data via conditional quantile correlations. <i>Journal of Multivariate Analysis</i> , 2018, 165, 1-13.	0.5	1
78	A Blockwise Consistency Method for Parameter Estimation of Complex Models. <i>Sankhya B</i> , 2018, 80, 179-223.	0.4	1
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80	Inverse Modeling: A Strategy to Cope with Non-linearity. <i>Springer Handbooks of Computational Statistics</i> , 2018, , 301-323.	0.2	0
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88	ExSIS: Extended sure independence screening for ultrahigh-dimensional linear models. <i>Signal Processing</i> , 2019, 159, 33-48.	2.1	4
89	Forward regression for Cox models with high-dimensional covariates. <i>Journal of Multivariate Analysis</i> , 2019, 173, 268-290.	0.5	11
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100	Conditional distance correlation screening for sparse ultrahigh-dimensional models. <i>Applied Mathematical Modelling</i> , 2020, 81, 232-252.	2.2	4
101	High-dimensional spectral data classification with nonparametric feature screening. <i>Journal of Chemometrics</i> , 2020, 34, e3199.	0.7	1
102	Nonparametric screening and feature selection for ultrahigh-dimensional Case II interval-censored failure time data. <i>Biometrical Journal</i> , 2020, 62, 1909-1925.	0.6	3
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104	Non-marginal feature screening for additive hazard model with ultrahigh-dimensional covariates. <i>Communications in Statistics - Theory and Methods</i> , 2020, , 1-19.	0.6	0
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113	Model-Free Feature Screening and FDR Control With Knockoff Features. <i>Journal of the American Statistical Association</i> , 2022, 117, 428-443.	1.8	26
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115	High-dimensional variable screening under multicollinearity. <i>Stat</i> , 2020, 9, e272.	0.3	9
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142	Focused Estimation and Model Averaging for High-Dimensional Data, An Overview. SSRN Electronic Journal, 0, , .	0.4	0
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157	Advanced OLS for Business Data Analytics. , 2021, , 279-312.		0
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159	Adaptive and powerful microbiome multivariate association analysis via feature selection. <i>NAR Genomics and Bioinformatics</i> , 2022, 4, lqab120.	1.5	3
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