List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Association of Delirium With Long-term Cognitive Decline. JAMA Neurology, 2020, 77, 1373.	4.5	226
2	Optimizing Treatment of Complicated Grief. JAMA Psychiatry, 2016, 73, 685.	6.0	222
3	Treatment of Complicated Grief in Elderly Persons. JAMA Psychiatry, 2014, 71, 1287.	6.0	206
4	Patient adherence predicts outcome from cognitive behavioral therapy in obsessive-compulsive disorder Journal of Consulting and Clinical Psychology, 2011, 79, 247-252.	1.6	176
5	Implementing Coordinated Specialty Care for Early Psychosis: The RAISE Connection Program. Psychiatric Services, 2015, 66, 691-698.	1.1	140
6	Age-specific penetrance of <i>LRRK2</i> G2019S in the Michael J. Fox Ashkenazi Jewish LRRK2 Consortium. Neurology, 2015, 85, 89-95.	1.5	130
7	Penetrance estimate of <i>LRRK2</i> p.G2019S mutation in individuals of nonâ€Ashkenazi Jewish ancestry. Movement Disorders, 2017, 32, 1432-1438.	2.2	126
8	Olanzapine Versus Placebo in Adult Outpatients With Anorexia Nervosa: A Randomized Clinical Trial. American Journal of Psychiatry, 2019, 176, 449-456.	4.0	116
9	Methods for Analyzing Multivariate Phenotypes in Genetic Association Studies. Journal of Probability and Statistics, 2012, 2012, 1-13.	0.3	90
10	Learning Optimal Personalized Treatment Rules in Consideration of Benefit and Risk: With an Application to Treating Type 2 Diabetes Patients With Insulin Therapies. Journal of the American Statistical Association, 2018, 113, 1-13.	1.8	66
11	Meta-analysis of mismatch negativity to simple versus complex deviants in schizophrenia. Schizophrenia Research, 2018, 191, 25-34.	1.1	64
12	Risk of Parkinson Disease in Carriers of Parkin Mutations. Archives of Neurology, 2008, 65, 467.	4.9	55
13	Complicated grief after suicide bereavement and other causes of death. Death Studies, 2017, 41, 267-275.	1.8	55
14	A Penalized Spline Approach to Functional Mixed Effects Model Analysis. Biometrics, 2011, 67, 861-870.	0.8	53
15	Augmented outcomeâ€weighted learning for estimating optimal dynamic treatment regimens. Statistics in Medicine, 2018, 37, 3776-3788.	0.8	41
16	Variable selection and prediction with incomplete high-dimensional data. Annals of Applied Statistics, 2016, 10, 418-450.	0.5	36
17	The impact of emotion awareness and regulation on psychotic symptoms during daily functioning. NPJ Schizophrenia, 2020, 6, 7.	2.0	32
18	Predictors of illness course and health maintenance following inpatient treatment among patients with anorexia nervosa. Journal of Eating Disorders, 2020, 8, 69.	1.3	31

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19	Inflammatory biomarkers in psychosis and clinical high risk populations. Schizophrenia Research, 2019, 206, 440-443.	1.1	30
20	Use of mobile assessment technologies in inpatient psychiatric settings. Asian Journal of Psychiatry, 2014, 10, 90-95.	0.9	29
21	Abnormal eating behavior in video-recorded meals in anorexia nervosa. Eating Behaviors, 2015, 19, 28-32.	1.1	28
22	Physical activity and postâ€treatment weight trajectory in anorexia nervosa. International Journal of Eating Disorders, 2016, 49, 482-489.	2.1	25
23	Measuring Maladaptive Cognitions in Complicated Grief: Introducing the Typical Beliefs Questionnaire. American Journal of Geriatric Psychiatry, 2017, 25, 541-550.	0.6	24
24	Efficient <i>â""</i> ₀ â€norm feature selection based on augmented and penalized minimization. Statistics in Medicine, 2018, 37, 473-486.	0.8	24
25	Crowdsourcing digital health measures to predict Parkinson's disease severity: the Parkinson's Disease Digital Biomarker DREAM Challenge. Npj Digital Medicine, 2021, 4, 53.	5.7	24
26	Treatment of Complicated Grief in Survivors of Suicide Loss. Journal of Clinical Psychiatry, 2018, 79, 17m11592.	1.1	24
27	Time Course of Relapse Following Acute Treatment for Anorexia Nervosa. American Journal of Psychiatry, 2021, 178, 848-853.	4.0	23
28	Assessment of test-retest reliability of a food choice task among healthy individuals. Appetite, 2018, 123, 352-356.	1.8	22
29	Longitudinal Association Analysis of Quantitative Traits. Genetic Epidemiology, 2012, 36, 856-869.	0.6	21
30	A Ridge Penalized Principal-Components Approach Based on Heritability for High-Dimensional Data. Human Heredity, 2007, 64, 182-191.	0.4	20
31	Combining isotonic regression and EM algorithm to predict genetic risk under monotonicity constraint. Annals of Applied Statistics, 2014, 8, 1182-1208.	0.5	19
32	Yearning and Its Measurement in Complicated Grief. Journal of Loss and Trauma, 2016, 21, 410-420.	0.9	19
33	Multivariate analysis of complex gene expression and clinical phenotypes with genetic marker data. Genetic Epidemiology, 2007, 31, S103-S109.	0.6	17
34	Nonparametric Estimation for Censored Mixture Data With Application to the Cooperative Huntington's Observational Research Trial. Journal of the American Statistical Association, 2012, 107, 1324-1338.	1.8	17
35	Prediction-Based Structured Variable Selection through the Receiver Operating Characteristic Curves. Biometrics, 2011, 67, 896-905.	0.8	16
36	Flexible Semiparametric Analysis of Longitudinal Genetic Studies by Reduced Rank Smoothing. Journal of the Royal Statistical Society Series C: Applied Statistics, 2012, 61, 1-24.	0.5	16

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37	Survival-Convolution Models for Predicting COVID-19 Cases and Assessing Effects of Mitigation Strategies. Frontiers in Public Health, 2020, 8, 325.	1.3	16
38	Transcription activity hot spot, is it real or an artifact?. BMC Proceedings, 2007, 1, S94.	1.8	12
39	A Marginal Approach to Reduced-Rank Penalized Spline Smoothing With Application to Multilevel Functional Data. Journal of the American Statistical Association, 2013, 108, 1216-1229.	1.8	12
40	Multiple Kernel Learning with Random Effects for Predicting Longitudinal Outcomes and Data Integration. Biometrics, 2015, 71, 918-928.	0.8	12
41	Cognitive training to improve memory in individuals undergoing electroconvulsive therapy: Negative findings. Journal of Psychiatric Research, 2017, 92, 8-14.	1.5	11
42	Estimation and Evaluation of Linear Individualized Treatment Rules to Guarantee Performance. Biometrics, 2018, 74, 517-528.	0.8	11
43	Maximizing remission from cognitive-behavioral therapy in medicated adults with obsessive-compulsive disorder. Behaviour Research and Therapy, 2021, 143, 103890.	1.6	11
44	A Method for Estimating Penetrance from Families Sampled for Linkage Analysis. Biometrics, 2006, 62, 1081-1088.	0.8	10
45	On Testing an Unspecified Function Through a Linear Mixed Effects Model with Multiple Variance Components. Biometrics, 2012, 68, 1113-1125.	0.8	10
46	Grief-related panic symptoms in Complicated Grief. Journal of Affective Disorders, 2015, 170, 213-216.	2.0	10
47	Matched Learning for Optimizing Individualized Treatment Strategies Using Electronic Health Records. Journal of the American Statistical Association, 2020, 115, 380-392.	1.8	10
48	Predicting Disease Onset from Mutation Status Using Proband and Relative Data with Applications to Huntington's Disease. Journal of Probability and Statistics, 2012, 2012, 1-19.	0.3	9
49	Efficient distribution estimation for data with unobserved sub-population identifiers. Electronic Journal of Statistics, 2012, 6, 710-737.	0.4	9
50	Examination of factor structure of the inventory of complicated grief (ICG) in a sample of bereaved military family members with persistent and elevated grief. International Journal of Methods in Psychiatric Research, 2017, 26, .	1.1	9
51	Sequential Multiple Assignment Randomization Trials with Enrichment Design. Biometrics, 2017, 73, 378-390.	0.8	9
52	Joint Rare Variant Association Test of the Average and Individual Effects for Sequencing Studies. PLoS ONE, 2012, 7, e32485.	1.1	9
53	Testing for familial aggregation of functional traits. Statistics in Medicine, 2009, 28, 3611-3625.	0.8	8
54	The reinforcing effect of exercise in anorexia nervosa: Clinical correlates and relationship to outcome. Eating Disorders, 2016, 24, 412-423.	1.9	8

YUANJIA WANG

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55	Clustering and principal-components approach based on heritability for mapping multiple gene expressions. BMC Proceedings, 2007, 1, S121.	1.8	7
56	Time-varying proportional odds model for mega-analysis of clustered event times. Biostatistics, 2019, 20, 129-146.	0.9	7
57	Analysis of genome-wide association data by large-scale Bayesian logistic regression. BMC Proceedings, 2009, 3, S16.	1.8	6
58	Estimating disease onset distribution functions in mutation carriers with censored mixture data. Journal of the Royal Statistical Society Series C: Applied Statistics, 2014, 63, 1-23.	0.5	6
59	Composite interaction tree for simultaneous learning of optimal individualized treatment rules and subgroups. Statistics in Medicine, 2019, 38, 2632-2651.	0.8	6
60	Identifying diseaseâ€associated biomarker network features through conditional graphical model. Biometrics, 2020, 76, 995-1006.	0.8	6
61	Impulsivity in cocaine users compared to matched controls: Effects of sex and preferred route of cocaine use. Drug and Alcohol Dependence, 2021, 226, 108840.	1.6	6
62	Auxiliary Marker-Assisted Classification in the Absence of Class Identifiers. Journal of the American Statistical Association, 2013, 108, 553-565.	1.8	5
63	Efficient estimation of nonparametric genetic risk function with censored data. Biometrika, 2015, 102, 515-532.	1.3	5
64	Penalized Nonlinear Mixed Effects Model to Identify Biomarkers That Predict Disease Progression. Biometrics, 2017, 73, 1343-1354.	0.8	5
65	Performance and psychometric properties of the Interpersonal Support Evaluation List (ISEL) in older adults with Complicated Grief. Journal of Affective Disorders, 2017, 218, 388-393.	2.0	5
66	Estimating personalized diagnostic rules depending on individualized characteristics. Statistics in Medicine, 2017, 36, 1099-1117.	0.8	5
67	Statistical modeling of Huntington disease onset. Handbook of Clinical Neurology / Edited By P J Vinken and G W Bruyn, 2017, 144, 47-61.	1.0	5
68	Outcome-Weighted Learning for Personalized Medicine with Multiple Treatment Options. , 2018, 2018, 565-574.		5
69	Modelling temporal biomarkers with semiparametric nonlinear dynamical systems. Biometrika, 2021, 108, 199-214.	1.3	5
70	Optimizing relapse prevention and changing habits (<scp>REACH</scp> +) in anorexia nervosa. International Journal of Eating Disorders, 2022, 55, 851-857.	2.1	5
71	Flexible estimation of covariance function by penalized spline with application to longitudinal family data. Statistics in Medicine, 2011, 30, 1883-1897.	0.8	4
72	Targeted Local Support Vector Machine for Age-Dependent Classification. Journal of the American Statistical Association, 2014, 109, 1174-1187.	1.8	4

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73	Learning Individualized Treatment Rules for Multiple-Domain Latent Outcomes. Journal of the American Statistical Association, 2021, 116, 269-282.	1.8	4
74	Age of Onset of Huntington's Disease in Carriers of Reduced Penetrance Alleles. Movement Disorders, 2021, 36, 2958-2961.	2.2	4
75	Evaluating effectiveness of public health intervention strategies for mitigating COVIDâ€19 pandemic. Statistics in Medicine, 2022, 41, 3820-3836.	0.8	4
76	Fused kernel-spline smoothing for repeatedly measured outcomes in a generalized partially linear model with functional single index. Annals of Statistics, 2015, 43, 1929-1958.	1.4	3
77	Estimation of genetic risk function with covariates in the presence of missing genotypes. Statistics in Medicine, 2017, 36, 3533-3546.	0.8	3
78	Nonlinear model with random inflection points for modeling neurodegenerative disease progression. Statistics in Medicine, 2018, 37, 4721-4742.	0.8	3
79	Use of personalized Dynamic Treatment Regimes (DTRs) and Sequential Multiple Assignment Randomized Trials (SMARTs) in mental health studies. Shanghai Archives of Psychiatry, 2014, 26, 376-83.	0.7	3
80	Unbiased and Locally Efficient Estimation of Genetic Effect on Quantitative Trait in the Presence of Population Admixture. Biometrics, 2011, 67, 331-343.	0.8	2
81	Semiparametric variance components models for genetic studies with longitudinal phenotypes. Biostatistics, 2012, 13, 482-496.	0.9	2
82	Nonparametric modeling and analysis of association between Huntington's disease onset and CAG repeats. Statistics in Medicine, 2014, 33, 1369-1382.	0.8	2
83	Learning Subject-Specific Directed Acyclic Graphs With Mixed Effects Structural Equation Models From Observational Data. Frontiers in Genetics, 2018, 9, 430.	1.1	2
84	On using electronic health records to improve optimal treatment rules in randomized trials. Biometrics, 2020, 76, 1075-1086.	0.8	2
85	Estimating disease onset from change points of markers measured with error. Biostatistics, 2021, 22, 819-835.	0.9	2
86	A General Framework of Nonparametric Feature Selection in High-Dimensional Data. Biometrics, 2023, 79, 951-963.	0.8	2
87	Predicting cumulative risk of disease onset by redistributing weights. Statistics in Medicine, 2015, 34, 2427-2443.	0.8	1
88	Learning Personalized Treatment Rules from Electronic Health Records Using Topic Modeling Feature Extraction. , 2019, 2019, 392-402.		1
89	Learning latent heterogeneity for type 2 diabetes patients using longitudinal health markers in electronic health records. Statistics in Medicine, 2021, 40, 1930-1946.	0.8	1
90	Integrative network learning for multimodality biomarker data. Annals of Applied Statistics, 2021, 15, 64-87.	0.5	1

YUANJIA WANG

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91	Semiparametric regression analysis of bivariate censored events in a family study of Alzheimer's disease. Biostatistics, 2022, 24, 32-51.	0.9	1
92	Statistical Learning Methods for Optimizing Dynamic Treatment Regimes in Subgroup Identification. Emerging Topics in Statistics and Biostatistics, 2020, , 271-297.	0.1	1
93	Heterogeneity of treatment effects. Shanghai Archives of Psychiatry, 2012, 24, 50-1.	0.7	1
94	Support Vector Hazards Machine: A Counting Process Framework for Learning Risk Scores for Censored Outcomes. Journal of Machine Learning Research, 2016, 17, .	62.4	1
95	Conditional Gaussian graphical model for estimating personalized disease symptom networks. Statistics in Medicine, 2022, 41, 543-553.	0.8	1
96	Efficient Nonparametric Estimation from Kin–Cohort Data. Communications in Statistics - Theory and Methods, 2010, 39, 3622-3634.	0.6	0
97	Assessment of a disease screener by hierarchical allâ€subset selection using area under the receiver operating characteristic curves. Statistics in Medicine, 2011, 30, 1751-1760.	0.8	0
98	A note on a nonparametric regression test through penalized splines. Statistica Sinica, 2014, 24, 1143-1160.	0.2	0
99	Leveraging Nonlinear Dynamic Models to Predict Progression of Neuroimaging Biomarkers. Biometrics, 2019, 75, 1240-1252.	0.8	0
100	Integrating multipleâ€domain rules for disease classification. Statistics in Medicine, 2019, 38, 3091-3104.	0.8	0
101	Constructing disease onset signatures using multi-dimensional network-structured biomarkers. Biostatistics, 2020, 21, 122-138.	0.9	0
102	Synthesizing independent stagewise trials for optimal dynamic treatment regimes. Statistics in Medicine, 2020, 39, 4107-4119.	0.8	0
103	Sparse nonparametric regression with regularized tensor product kernel. Stat, 2020, 9, .	0.3	0
104	Time-varying Hazards Model for Incorporating Irregularly Measured, High-Dimensional Biomarkers. Statistica Sinica, 2020, 30, 1605-1632.	0.2	0
105	Selfâ€matched learning to construct treatment decision rulesÂfrom electronic health records. Statistics in Medicine, 2022, , .	0.8	0