

Ruoqing Zhu

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2755604/publications.pdf>

Version: 2024-02-01

25
papers

630
citations

687363

13
h-index

677142

22
g-index

26
all docs

26
docs citations

26
times ranked

841
citing authors

#	ARTICLE	IF	CITATIONS
1	Efficient gradient boosting for prognostic biomarker discovery. <i>Bioinformatics</i> , 2022, 38, 1631-1638.	4.1	13
2	Dimension Reduction Forests: Local Variable Importance Using Structured Random Forests. <i>Journal of Computational and Graphical Statistics</i> , 2022, 31, 1104-1113.	1.7	1
3	Fecal Bacteria as Biomarkers for Predicting Food Intake in Healthy Adults. <i>Journal of Nutrition</i> , 2021, 151, 423-433.	2.9	26
4	Topic Modeling on Triage Notes With Semiorthogonal Nonnegative Matrix Factorization. <i>Journal of the American Statistical Association</i> , 2021, 116, 1609-1624.	3.1	5
5	Diagnostic and prognostic capabilities of a biomarker and EMR-based machine learning algorithm for sepsis. <i>Clinical and Translational Science</i> , 2021, 14, 1578-1589.	3.1	12
6	A parsimonious personalized dose-finding model via dimension reduction. <i>Biometrika</i> , 2021, 108, 643-659.	2.4	2
7	Lung epithelial and endothelial damage, loss of tissue repair, inhibition of fibrinolysis, and cellular senescence in fatal COVID-19. <i>Science Translational Medicine</i> , 2021, 13, eabj7790.	12.4	133
8	Risk assessment of latent tuberculosis infection through a multiplexed cytokine biosensor assay and machine learning feature selection. <i>Scientific Reports</i> , 2021, 11, 20544.	3.3	20
9	Nonparametric variable selection and its application to additive models. <i>Annals of the Institute of Statistical Mathematics</i> , 2020, 72, 827-854.	0.8	1
10	Assessment of peritoneal microbial features and tumor marker levels as potential diagnostic tools for ovarian cancer. <i>PLoS ONE</i> , 2020, 15, e0227707.	2.5	28
11	Constructing dynamic treatment regimes with shared parameters for censored data. <i>Statistics in Medicine</i> , 2020, 39, 1250-1263.	1.6	7
12	Differential Effects of Influenza Virus NA, HA Head, and HA Stalk Antibodies on Peripheral Blood Leukocyte Gene Expression during Human Infection. <i>MBio</i> , 2019, 10, .	4.1	8
13	Counting process-based dimension reduction methods for censored outcomes. <i>Biometrika</i> , 2019, 106, 181-196.	2.4	6
14	Bagging and Deep Learning in Optimal Individualized Treatment Rules. <i>Biometrics</i> , 2019, 75, 674-684.	1.4	22
15	GradientScanSurv—An exhaustive association test method for gene expression data with censored survival outcome. <i>PLoS ONE</i> , 2018, 13, e0207590.	2.5	2
16	Combining Biomarkers with EMR Data to Identify Patients in Different Phases of Sepsis. <i>Scientific Reports</i> , 2017, 7, 10800.	3.3	59
17	Greedy Outcome Weighted Tree Learning of Optimal Personalized Treatment Rules. <i>Biometrics</i> , 2017, 73, 391-400.	1.4	33
18	Tree based weighted learning for estimating individualized treatment rules with censored data. <i>Electronic Journal of Statistics</i> , 2017, 11, 3927-3953.	0.7	28

#	ARTICLE	IF	CITATIONS
19	Increasing Access to State Psychiatric Hospital Beds: Exploring Supply-Side Solutions. <i>Psychiatric Services</i> , 2016, 67, 523-528.	2.0	23
20	Integrating multidimensional omics data for cancer outcome. <i>Biostatistics</i> , 2016, 17, 605-618.	1.5	32
21	Reinforcement Learning Trees. <i>Journal of the American Statistical Association</i> , 2015, 110, 1770-1784.	3.1	105
22	Identifying Gene-Environment and Gene-Gene Interactions Using a Progressive Penalization Approach. <i>Genetic Epidemiology</i> , 2014, 38, 353-368.	1.3	16
23	Recursively Imputed Survival Trees. <i>Journal of the American Statistical Association</i> , 2012, 107, 331-340.	3.1	45
24	Estimating Heterogeneous Treatment Effect on Multivariate Responses Using Random Forests. <i>Statistics in Biosciences</i> , 0, , 1.	1.2	1
25	Dermoscopic Image Classification with Neural Style Transfer. <i>Journal of Computational and Graphical Statistics</i> , 0, , 1-30.	1.7	2