

Vera DjordjiloviÄ

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6066226/publications.pdf>

Version: 2024-02-01

14
papers

124
citations

1937457

4
h-index

1372474

10
g-index

14
all docs

14
docs citations

14
times ranked

228
citing authors

#	ARTICLE	IF	CITATIONS
1	The role of the frontal aslant tract and premotor connections in visually guided hand movements. <i>NeuroImage</i> , 2017, 146, 419-428.	2.1	50
2	Global test for high-dimensional mediation: Testing groups of potential mediators. <i>Statistics in Medicine</i> , 2019, 38, 3346-3360.	0.8	26
3	High-throughput mediation analysis of human proteome and metabolome identifies mediators of post-bariatric surgical diabetes control. <i>Nature Communications</i> , 2021, 12, 6951.	5.8	13
4	Leveraging three-dimensional chromatin architecture for effective reconstruction of enhancer-target gene regulatory interactions. <i>Nucleic Acids Research</i> , 2021, 49, e97-e97.	6.5	6
5	Multi split conformal prediction. <i>Statistics and Probability Letters</i> , 2022, 184, 109395.	0.4	5
6	On optimal two-stage testing of multiple mediators. <i>Biometrical Journal</i> , 2022, , .	0.6	5
7	Graphical modeling for gene set analysis: A critical appraisal. <i>Biometrical Journal</i> , 2015, 57, 852-866.	0.6	4
8	SourceSet: A graphical model approach to identify primary genes in perturbed biological pathways. <i>PLoS Computational Biology</i> , 2019, 15, e1007357.	1.5	4
9	Lifetime Ultraviolet Radiation Exposure and DNA Methylation in Blood Leukocytes: The Norwegian Women and Cancer Study. <i>Scientific Reports</i> , 2020, 10, 4521.	1.6	4
10	simPATHY: a new method for simulating data from perturbed biological PATHways. <i>Bioinformatics</i> , 2017, 33, 456-457.	1.8	3
11	An empirical comparison of popular structure learning algorithms with a view to gene network inference. <i>International Journal of Approximate Reasoning</i> , 2017, 88, 602-613.	1.9	3
12	Searching for a source of difference in graphical models. <i>Journal of Multivariate Analysis</i> , 2022, 190, 104973.	0.5	1
13	Simulating gene silencing through intervention analysis. <i>Journal of the Royal Statistical Society Series C: Applied Statistics</i> , 2020, 69, 887-907.	0.5	0
14	Searching for the Source of Difference: A Graphical Model Approach. <i>Lecture Notes in Computer Science</i> , 2020, , 131-138.	1.0	0