

MarÃ-a MartÃ-nez Ballesteros

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8694063/publications.pdf>

Version: 2024-02-01

23
papers

410
citations

687363

13
h-index

752698

20
g-index

24
all docs

24
docs citations

24
times ranked

382
citing authors

#	ARTICLE	IF	CITATIONS
1	Mining quantitative association rules based on evolutionary computation and its application to atmospheric pollution. <i>Integrated Computer-Aided Engineering</i> , 2010, 17, 227-242.	4.6	49
2	An evolutionary algorithm to discover quantitative association rules in multidimensional time series. <i>Soft Computing</i> , 2011, 15, 2065-2084.	3.6	42
3	MRQAR: A generic MapReduce framework to discover quantitative association rules in big data problems. <i>Knowledge-Based Systems</i> , 2018, 153, 176-192.	7.1	35
4	Selecting the best measures to discover quantitative association rules. <i>Neurocomputing</i> , 2014, 126, 3-14.	5.9	32
5	Enhancing the scalability of a genetic algorithm to discover quantitative association rules in large-scale datasets. <i>Integrated Computer-Aided Engineering</i> , 2015, 22, 21-39.	4.6	31
6	Autoencoded DNA methylation data to predict breast cancer recurrence: Machine learning models and gene-weight significance. <i>Artificial Intelligence in Medicine</i> , 2020, 110, 101976.	6.5	27
7	External clustering validity index based on chi-squared statistical test. <i>Information Sciences</i> , 2019, 487, 1-17.	6.9	26
8	Discovering gene association networks by multi-objective evolutionary quantitative association rules. <i>Journal of Computer and System Sciences</i> , 2014, 80, 118-136.	1.2	25
9	Machine learning techniques to discover genes with potential prognosis role in Alzheimer's disease using different biological sources. <i>Information Fusion</i> , 2017, 36, 114-129.	19.1	22
10	A Nearest Neighbours-Based Algorithm for Big Time Series Data Forecasting. <i>Lecture Notes in Computer Science</i> , 2016, , 174-185.	1.3	20
11	An approach to validity indices for clustering techniques in Big Data. <i>Progress in Artificial Intelligence</i> , 2018, 7, 81-94.	2.4	19
12	Improving a multi-objective evolutionary algorithm to discover quantitative association rules. <i>Knowledge and Information Systems</i> , 2016, 49, 481-509.	3.2	15
13	A study of the suitability of autoencoders for preprocessing data in breast cancer experimentation. <i>Journal of Biomedical Informatics</i> , 2017, 72, 33-44.	4.3	15
14	Evolutionary association rules for total ozone content modeling from satellite observations. <i>Chemometrics and Intelligent Laboratory Systems</i> , 2011, 109, 217-227.	3.5	12
15	Obtaining optimal quality measures for quantitative association rules. <i>Neurocomputing</i> , 2016, 176, 36-47.	5.9	11
16	Analysis of Measures of Quantitative Association Rules. <i>Lecture Notes in Computer Science</i> , 2011, , 319-326.	1.3	7
17	Inferring gene-gene associations from Quantitative Association Rules. , 2011, , .		6
18	Analysis of the Evolution of the Spanish Labour Market Through Unsupervised Learning. <i>IEEE Access</i> , 2019, 7, 121695-121708.	4.2	6

#	ARTICLE	IF	CITATIONS
19	Quantitative Association Rules Applied to Climatological Time Series Forecasting. Lecture Notes in Computer Science, 2009, , 284-291.	1.3	6
20	A Sensitivity Analysis for Quality Measures of Quantitative Association Rules. Lecture Notes in Computer Science, 2013, , 578-587.	1.3	2
21	Applications of Computational Intelligence in Time Series. Computational Intelligence and Neuroscience, 2017, 2017, 1-2.	1.7	1
22	On the use of algorithms to discover motifs in DNA sequences. , 2011, , .		0
23	A novel approach to discover numerical association based on the coronavirus optimization algorithm. , 2022, , .		0