Francisco Gmez-Vela

List of Publications by Citations

Source: https://exaly.com/author-pdf/1137752/francisco-gomez-vela-publications-by-citations.pdf

Version: 2024-04-19

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

34 381 10 19 g-index

43 555 avg, IF 4.23 L-index

#	Paper	IF	Citations
34	High-dimensional feature selection via feature grouping: A Variable Neighborhood Search approach. <i>Information Sciences</i> , 2016 , 326, 102-118	7.7	66
33	Stacking Ensemble Learning for Short-Term Electricity Consumption Forecasting. <i>Energies</i> , 2018 , 11, 949	3.1	63
32	Computational methods for Gene Regulatory Networks reconstruction and analysis: A review. <i>Artificial Intelligence in Medicine</i> , 2019 , 95, 133-145	7.4	57
31	A Comparative Study of Time Series Forecasting Methods for Short Term Electric Energy Consumption Prediction in Smart Buildings. <i>Energies</i> , 2019 , 12, 1934	3.1	30
30	An effective measure for assessing the quality of biclusters. <i>Computers in Biology and Medicine</i> , 2012 , 42, 245-56	7	25
29	Social symbol grounding and language evolution. <i>Interaction Studies</i> , 2007 , 8, 31-52	1.3	23
28	Pangenome of uncovers two groups of genomes, one of them with genes involved in CRISPR/Cas defence systems associated with the absence of plasmids and exclusive genes for biofilm formation. <i>Microbial Genomics</i> , 2019 , 5,	4.4	21
27	Gene network coherence based on prior knowledge using direct and indirect relationships. <i>Computational Biology and Chemistry</i> , 2015 , 56, 142-51	3.6	11
26	A multivariate approach to the symmetrical uncertainty measure: Application to feature selection problem. <i>Information Sciences</i> , 2019 , 494, 1-20	7.7	10
25	Identifying livestock behavior patterns based on accelerometer dataset. <i>Journal of Computational Science</i> , 2020 , 41, 101076	3.4	10
24	Incorporating biological knowledge for construction of fuzzy networks of gene associations. <i>Applied Soft Computing Journal</i> , 2016 , 42, 144-155	7.5	9
23	GNC-app: A new Cytoscape app to rate gene networks biological coherence using gene-gene indirect relationships. <i>BioSystems</i> , 2018 , 166, 61-65	1.9	8
22	Hybridizing Deep Learning and Neuroevolution: Application to the Spanish Short-Term Electric Energy Consumption Forecasting. <i>Applied Sciences (Switzerland)</i> , 2020 , 10, 5487	2.6	7
21	GFD-Net: A novel semantic similarity methodology for the analysis of gene networks. <i>Journal of Biomedical Informatics</i> , 2017 , 68, 71-82	10.2	6
20	Structure Optimization for Large Gene Networks Based on Greedy Strategy. <i>Computational and Mathematical Methods in Medicine</i> , 2018 , 2018, 9674108	2.8	6
19	Gene network biological validity based on gene-gene interaction relevance. <i>Scientific World Journal, The</i> , 2014 , 2014, 540679	2.2	5
18	Pattern Recognition in Biological Time Series. <i>Lecture Notes in Computer Science</i> , 2011 , 164-172	0.9	4

LIST OF PUBLICATIONS

17	A Comparative Study of Supervised Machine Learning Algorithms for the Prediction of Long-Range Chromatin Interactions. <i>Genes</i> , 2020 , 11,	4.2	4	
16	Computational Inference of Gene Co-Expression Networks for the identification of Lung Carcinoma Biomarkers: An Ensemble Approach. <i>Genes</i> , 2019 , 10,	4.2	3	
15	Genome-wide prediction of topoisomerase III binding by architectural factors and chromatin accessibility. <i>PLoS Computational Biology</i> , 2021 , 17, e1007814	5	3	
14	Biclustering of Smart Building Electric Energy Consumption Data. <i>Applied Sciences (Switzerland)</i> , 2019 , 9, 222	2.6	2	
13	Computational Analysis of the Global Effects of in the Immune Response to Coronavirus Infection Using Gene Networks. <i>Genes</i> , 2020 , 11,	4.2	2	
12	BIGO: A web application to analyse gene enrichment analysis results. <i>Computational Biology and Chemistry</i> , 2018 , 76, 169-178	3.6	1	
11	A multi-objective genetic algorithm for the Protein Structure Prediction 2011,		1	
10	Distribution level electric current consumption and meteorological data set of the east region of Paraguay <i>Data in Brief</i> , 2022 , 40, 107699	1.2	1	
9	Ensemble and Greedy Approach for the Reconstruction of Large Gene Co-Expression Networks. <i>Entropy</i> , 2019 , 21, 1139	2.8	1	
8	Analysis of Student Achievement Scores: A Machine Learning Approach. <i>Advances in Intelligent Systems and Computing</i> , 2020 , 275-284	0.4	1	
7	Analysis of Electric Energy Consumption Profiles Using a Machine Learning Approach: A Paraguayan Case Study. <i>Electronics (Switzerland)</i> , 2022 , 11, 267	2.6	O	
6	Gene Regulatory Networks Validation Framework Based in KEGG. <i>Lecture Notes in Computer Science</i> , 2011 , 279-286	0.9	O	
5	Automatic Diagnosis of Ocular Toxoplasmosis from Fundus Images with Residual Neural Networks. <i>Studies in Health Technology and Informatics</i> , 2021 , 281, 173-177	0.5	O	
4	Bioinformatics from a Big Data Perspective: Meeting the Challenge. <i>Lecture Notes in Computer Science</i> , 2017 , 349-359	0.9		
3	gMSR: A Multi-GPU Algorithm to Accelerate a Massive Validation of Biclusters. <i>Electronics</i> (Switzerland), 2020 , 9, 1782	2.6		
2	Redundancy Is Not Necessarily Detrimental in Classification Problems. <i>Mathematics</i> , 2021 , 9, 2899	2.3		
1	Analysis of Relevance and Redundance on Topoisomerase 2b (TOP2B) Binding Sites: A Feature Selection Approach. <i>Lecture Notes in Computer Science</i> , 2018 , 86-101	0.9		