

Habib Motieghader

List of Publications by Citations

Source: <https://exaly.com/author-pdf/1883642/habib-motieghader-publications-by-citations.pdf>

Version: 2024-04-23

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.

14
papers

198
citations

8
h-index

14
g-index

18
ext. papers

292
ext. citations

4
avg, IF

3.85
L-index

#	Paper	IF	Citations
14	A hybrid gene selection algorithm for microarray cancer classification using genetic algorithm and learning automata. <i>Informatics in Medicine Unlocked</i> , 2017 , 9, 246-254	5.3	58
13	FeatureSelect: a software for feature selection based on machine learning approaches. <i>BMC Bioinformatics</i> , 2019 , 20, 170	3.6	44
12	mRNA-miRNA bipartite network reconstruction to predict prognostic module biomarkers in colorectal cancer stage differentiation. <i>Molecular BioSystems</i> , 2017 , 13, 2168-2180		21
11	Gene biomarker discovery at different stages of Alzheimer using gene co-expression network approach. <i>Scientific Reports</i> , 2020 , 10, 12210	4.9	17
10	World Competitive Contests (WCC) algorithm: A novel intelligent optimization algorithm for biological and non-biological problems. <i>Informatics in Medicine Unlocked</i> , 2016 , 3, 15-28	5.3	13
9	Gene co-expression network approach for predicting prognostic microRNA biomarkers in different subtypes of breast cancer. <i>Genomics</i> , 2020 , 112, 135-143	4.3	11
8	mRNA and microRNA selection for breast cancer molecular subtype stratification using meta-heuristic based algorithms. <i>Genomics</i> , 2020 , 112, 3207-3217	4.3	10
7	Repurposing novel therapeutic candidate drugs for coronavirus disease-19 based on protein-protein interaction network analysis. <i>BMC Biotechnology</i> , 2021 , 21, 22	3.5	10
6	Sequential and Mixed Genetic Algorithm and Learning Automata (SGALA, MGALA) for Feature Selection in QSAR. <i>Iranian Journal of Pharmaceutical Research</i> , 2017 , 16, 533-553	1.1	7
5	A machine learning method based on the genetic and world competitive contests algorithms for selecting genes or features in biological applications. <i>Scientific Reports</i> , 2021 , 11, 3349	4.9	4
4	Drug repurposing for coronavirus (SARS-CoV-2) based on gene co-expression network analysis. <i>Scientific Reports</i> , 2021 , 11, 21872	4.9	2
3	LncRNA DLGAP1-AS2 overexpression associates with gastric tumorigenesis: a promising diagnostic and therapeutic target.. <i>Molecular Biology Reports</i> , 2022 , 1	2.8	0
2	Drug Repurposing for Alzheimer's Disease Based on Protein-Protein Interaction Network. <i>BioMed Research International</i> , 2021 , 2021, 1280237	3	0
1	mRNA-miRNA bipartite networks reconstruction in different tissues of bladder cancer based on gene co-expression network analysis.. <i>Scientific Reports</i> , 2022 , 12, 5885	4.9	0