Habib Motieghader

List of Publications by Year in descending order

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932766 996533 16 401 10 15 citations g-index h-index papers 18 18 18 336 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	A hybrid gene selection algorithm for microarray cancer classification using genetic algorithm and learning automata. Informatics in Medicine Unlocked, 2017, 9, 246-254.	1.9	86
2	FeatureSelect: a software for feature selection based on machine learning approaches. BMC Bioinformatics, 2019, 20, 170.	1.2	83
3	Gene biomarker discovery at different stages of Alzheimer using gene co-expression network approach. Scientific Reports, 2020, 10, 12210.	1.6	39
4	mRNA–miRNA bipartite network reconstruction to predict prognostic module biomarkers in colorectal cancer stage differentiation. Molecular BioSystems, 2017, 13, 2168-2180.	2.9	33
5	Repurposing novel therapeutic candidate drugs for coronavirus disease-19 based on protein-protein interaction network analysis. BMC Biotechnology, 2021, 21, 22.	1.7	29
6	World Competitive Contests (WCC) algorithm: A novel intelligent optimization algorithm for biological and non-biological problems. Informatics in Medicine Unlocked, 2016, 3, 15-28.	1.9	25
7	Gene co-expression network approach for predicting prognostic microRNA biomarkers in different subtypes of breast cancer. Genomics, 2020, 112, 135-143.	1.3	22
8	mRNA and microRNA selection for breast cancer molecular subtype stratification using meta-heuristic based algorithms. Genomics, 2020, 112, 3207-3217.	1.3	18
9	A machine learning method based on the genetic and world competitive contests algorithms for selecting genes or features in biological applications. Scientific Reports, 2021, 11, 3349.	1.6	13
10	Drug repurposing for coronavirus (SARS-CoV-2) based on gene co-expression network analysis. Scientific Reports, 2021, 11, 21872.	1.6	13
11	Drug Repurposing for Alzheimer's Disease Based on Protein-Protein Interaction Network. BioMed Research International, 2021, 2021, 1-11.	0.9	11
12	Drug repositioning in non-small cell lung cancer (NSCLC) using gene co-expression and drug–gene interaction networks analysis. Scientific Reports, 2022, 12, .	1.6	10
13	Sequential and Mixed Genetic Algorithm and Learning Automata (SGALA, MGALA) for Feature Selection in QSAR. Iranian Journal of Pharmaceutical Research, 2017, 16, 533-553.	0.3	7
14	mRNA–miRNA bipartite networks reconstruction in different tissues of bladder cancer based on gene co-expression network analysis. Scientific Reports, 2022, 12, 5885.	1.6	6
15	LncRNA DLGAP1-AS2 overexpression associates with gastric tumorigenesis: a promising diagnostic and therapeutic target. Molecular Biology Reports, 2022, 49, 6817-6826.	1.0	5
16	Identification of Novel MicroRNAs and Their Targets in Leukemia Cancers: A Computational Approach. The Cancer Press, 2016, 1, 14.	0.1	0