Habib Motieghader

List of Publications by Year in Descending Order

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