Yosef Masoudi-sobhanzadeh

List of Publications by Year in Descending Order

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

15	195	7	13
papers	citations	h-index	g-index
16	294	5.3	4.14
ext. papers	ext. citations	avg, IF	L-index

#	Paper	IF	Citations
15	Pharmacoinformatics-based phytochemical screening for anticancer impacts of yellow sweet clover, Melilotus officinalis (Linn.) Pall. <i>Computers in Biology and Medicine</i> , 2021 , 138, 104921	7	2
14	Structure-based drug repurposing against COVID-19 and emerging infectious diseases: methods, resources and discoveries. <i>Briefings in Bioinformatics</i> , 2021 , 22,	13.4	2
13	Advances in the computational landscape for repurposed drugs against COVID-19. <i>Drug Discovery Today</i> , 2021 , 26, 2800-2815	8.8	4
12	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
11	A novel multi-objective metaheuristic algorithm for protein-peptide docking and benchmarking on the LEADS-PEP dataset. <i>Computers in Biology and Medicine</i> , 2021 , 138, 104896	7	3
10	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
9	Computational-based drug repurposing methods in COVID-19. <i>BioImpacts</i> , 2020 , 10, 205-206	3.5	9
8	Synthetic repurposing of drugs against hypertension: a datamining method based on association rules and a novel discrete algorithm. <i>BMC Bioinformatics</i> , 2020 , 21, 313	3.6	6
7	Drug databases and their contributions to drug repurposing. <i>Genomics</i> , 2020 , 112, 1087-1095	4.3	38
6	DrugR+: A comprehensive relational database for drug repurposing, combination therapy, and replacement therapy. <i>Computers in Biology and Medicine</i> , 2019 , 109, 254-262	7	21
5	FeatureSelect: a software for feature selection based on machine learning approaches. <i>BMC Bioinformatics</i> , 2019 , 20, 170	3.6	44
4	Trader as a new optimization algorithm predicts drug-target interactions efficiently. <i>Scientific Reports</i> , 2019 , 9, 9348	4.9	27
3	Disease global behavior: A systematic study of the human interactome network reveals conserved topological features among categories of diseases. <i>Informatics in Medicine Unlocked</i> , 2019 , 17, 100249	5.3	5
2	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
1	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