

# Hao Wu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9607266/publications.pdf>

Version: 2024-02-01

9  
papers

301  
citations

1307594

7  
h-index

1474206

9  
g-index

10  
all docs

10  
docs citations

10  
times ranked

424  
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrating Protein-Protein Interaction Networks and Somatic Mutation Data to Detect Driver Modules in Pan-Cancer. <i>Interdisciplinary Sciences, Computational Life Sciences</i> , 2022, 14, 151-167.	3.6	4
2	scHiCStackL: a stacking ensemble learning-based method for single-cell Hi-C classification using cell embedding. <i>Briefings in Bioinformatics</i> , 2022, 23, .	6.5	16
3	StackTADB: a stacking-based ensemble learning model for predicting the boundaries of topologically associating domains (TADs) accurately in fruit flies. <i>Briefings in Bioinformatics</i> , 2022, 23, .	6.5	12
4	Large-scale comparative review and assessment of computational methods for anti-cancer peptide identification. <i>Briefings in Bioinformatics</i> , 2021, 22, .	6.5	40
5	Strategies of attack-defense game for wireless sensor networks considering the effect of confidence level in fuzzy environment. <i>Engineering Applications of Artificial Intelligence</i> , 2021, 102, 104238.	8.1	10
6	Network-based method for detecting dysregulated pathways in glioblastoma cancer. <i>IET Systems Biology</i> , 2018, 12, 39-44.	1.5	5
7	Identifying overlapping mutated driver pathways by constructing gene networks in cancer. <i>BMC Bioinformatics</i> , 2015, 16, S3.	2.6	21
8	Detecting Overlapping Protein Complexes by Rough-Fuzzy Clustering in Protein-Protein Interaction Networks. <i>PLoS ONE</i> , 2014, 9, e91856.	2.5	43
9	A Network Based Method for Analysis of lncRNA-Disease Associations and Prediction of lncRNAs Implicated in Diseases. <i>PLoS ONE</i> , 2014, 9, e87797.	2.5	150