Hao Wu

List of Publications by Year in descending order

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1307594 1474206 9 301 7 9 citations g-index h-index papers 10 10 10 424 citing authors docs citations times ranked all docs

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