

Yang Liu

List of Publications by Year in descending order

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Version: 2024-02-01

8
papers

703
citations

1307594
7
h-index

1588992
8
g-index

9
all docs

9
docs citations

9
times ranked

412
citing authors

| # | ARTICLE | IF | CITATIONS |
|---|--|------|-----------|
| 1 | CB-Dock: a web server for cavity detection-guided protein–ligand blind docking. <i>Acta Pharmacologica Sinica</i> , 2020, 41, 138-144. | 6.1 | 377 |
| 2 | CB-Dock2: improved protein–ligand blind docking by integrating cavity detection, docking and homologous template fitting. <i>Nucleic Acids Research</i> , 2022, 50, W159-W164. | 14.5 | 219 |
| 3 | FitDock: protein–ligand docking by template fitting. <i>Briefings in Bioinformatics</i> , 2022, 23, . | 6.5 | 38 |
| 4 | FBXL2 counteracts Grp94 to destabilize EGFR and inhibit EGFR-driven NSCLC growth. <i>Nature Communications</i> , 2021, 12, 5919. | 12.8 | 29 |
| 5 | A systematic analysis of miRNA markers and classification algorithms for forensic body fluid identification. <i>Briefings in Bioinformatics</i> , 2021, 22, . | 6.5 | 13 |
| 6 | DrugDevCovid19: An Atlas of Anti-COVID-19 Compounds Derived by Computer-Aided Drug Design. <i>Molecules</i> , 2022, 27, 683. | 3.8 | 11 |
| 7 | LigMate: A Multifeature Integration Algorithm for Ligand-Similarity-Based Virtual Screening. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 6044-6053. | 5.4 | 8 |
| 8 | A Review on the Methods of Peptide-MHC Binding Prediction. <i>Current Bioinformatics</i> , 2021, 15, 878-888. | 1.5 | 8 |